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Q95Y35; O1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
HYPOTHETICAL 38.2 KDA PROTEIN.
Y108G3AL.2.
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investigating biology. The C.
Science 282:2012-2018(1998).
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Hypothetical protein,
SEQUENCE 327 AA; 38186 MW;
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
STRAIN-C57BL/GJ; TISSUE=TESTIS;
MREDLINE-21085660; PubMed=11217851;
Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
Saito T., Okazaki Y., Gojobori T., Sono H., Kasukawa T., Saito R.,
Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
Kuehl P., Lewis S., Matsuo Y., Nikaio I., Pesole G., Quackenbush
Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio
Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
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01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
1110021E09RIK PROTEIN (FRAGMENT).
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"The sequence
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Mammalia; Eutheria;
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                                                                                                                                       TISSUE-WHOLE KERNELS;
Lawrence C.J., Malmberg R.L., Muszynski M.G., Dawe "Maximum likelihood methods reveal conservation of closely related kinesin families.";
J. Mol. Evol. 0:0-0(2001).
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
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Q93XF8;
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EMBL: AK015938; BAB30042.1; ...
MGD; MGI:1914171; 1110021E09Rik.
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NCBI_TaxID=4577;
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Panicoideae; Andropogoneae; Ze
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yta; Liliopsida; Poales; Poaceae; PACC cl
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ALIGNMENTS

AAW79207 RESULT

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AAW79207 standard;

Protein;

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Framework 3 region of human V kappa gene HUM5400

21-DEC-1998 AAW79207;

(first entry)

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05-MAR-1993;
09-SEP-1993;
29-MAR-1995;
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Inhibition of T-cell mediated immune response with anti-CD2
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93US-0027008.
93US-0119032.
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RESULT AABB A298 ID AABB A298 AC AABB XX AABB AA
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     This invention describes novel antibodies (Ab) in which the variable region (VR) of at least one chain and/or the VR of at least one heavy chain includes at least one of 7 specified sequences, or fragments of these sequences, or contain at least one light chain and/or heavy chain encoded by specific nucleic acid sequences (I) and (II), reproduced, or their fragments. The products of the invention have
                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                            Claim
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                                                                                                                                                                                                                              diagnosis
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and plasma cell
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                                            30-JUN-1994;
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                                                                                                                                         06-MAR-1996
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                                                                                        27-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    analysis for T-lymphocyte antigenic sequences using a computer algorithm. Residues not within a complementarity determining region, canonical structure or Vernier zone may be modified to reduce immunogenicity in humans (e.g. in sequence AAR92995). This method, which involves the introduction of only a few point mutations into T-cell epitope coding regions, is generally applicable in humanisation of mouse antibodies. The resulting humanised antibodies may be used e.g. as antitumour agents. They retain the antigen recognition of the
                                                                                                                                            Region
  12-OCT-1995
                                                 W09527045-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pattern
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-MAY-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR80082;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR80082 standard; Protein; 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g. as antitumour agents. They retain the antigen r original antibody, but are not immunogenic in humans.
                                                                                                                                                                                                Region
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                                                                                                                                                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                                                                                                                  Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Light chain; RT3; murine; catalytic antibody; bacteriophage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse derived light chain RT3 phage antibody pattern I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      produced by a mouse hybridoma. The sequence is partial, complementarity determining regions are omitted. The sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence represents residues from a human immunoglobulin with homology to the light chain variable region from monoclonal antibody non-logy (AAR92993), specific for epidermal growth factor receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying interspecies differences in amino acid sequence of Ig T-cell epitopes - by sequence comparison, also humanised antibodies contg. altered T-cell epitopes, retaining antigen specificity but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 14; Fig 3; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         not immunogenicity, esp. for tumour treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-130770/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 FTLKISR 7
|||||||
53 ftlkisr 59
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/note=
83..89
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43..49
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29..42
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15..28
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                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                                                                                                                                                                                                   "framework region
                                                                                                                                                                                                                                                                                                                  "complementarity determining
                                                                                                                                                                                                                                                                                                                                                                      "framework region
                                                                                                                                                                                                                  "complementarity determining
                                                                                             "complementarity determining N-terminal fragment"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 33;
Pred. No.
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Best Local
                                                      Treatment and preventive agent for hypercalcaemia - contg. one of anti-human para-thyroid-hormone-related protein monoclonal antibod a rodent or chimera monoclonal antibody, fused gene and cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT04629 encodes AAR80082 mouse derived light chain RT3 phage antibody. The DNA was used in the prepn. of catalytic antibody (CA) producing bacteriophage. The CAs can be used to activate/deactivate a biological function in an animal by enhancing the rate of cleavage, or formation of a specific bond within a mol. in vivo.
                                                                                                                                        15-MAY-1990;
                                                                                                                                                            15-MAY-1991;
                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                         Antihuman parathyroid hormone-related protein; monoclonal antibody; variable region; rodent/human chimeric MAb; constant region; PTHrP.
                                                                                                                                                                                                                                                                        Hypercalcaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Production of catalytic antibodies displayed on phage generating a gene library of antibody-derived domains it in phage vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chiswell D,
Martin MT,
                                                                                                WPI; 1992-320987/39.
                                                                                                                    (KANF ) KANEKA CORP.
                                                                                                                                                                                18-AUG-1992.
                                                                                                                                                                                                   JP04228089-A.
                                                                                                                                                                                                                                                                                           20-MAY-1998
                                                                                                                                                                                                                                                                                                                AAR27009;
                                                                                                                                                                                                                                                                                                                                   AAR27009 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT04629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             68
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                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA;
                                                                                                                                        90JP-0124581
                                                                                                                                                            91JP-0110565
                                                                                                                                                                                                                                                                       agent portion
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                                                                                                                                                                                                                                                                                                                                  peptide;
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Smith RG, Titmas RC, W
                                                                                                                                                                                                                                                                                                                                   92
                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                          Score 33; DE
Pred. No. 3.6
0; Mismatches
                                                                                                                                                                                                                                                                                                                                   Å
                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 3.6;
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                                                                 antibody,
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The sequences given in AAR27008-11 are fragments which were used the construction of an agent for the treating and prevention of

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Disclosure; Page 13; 18pp; Japanese

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                                                   Query Match
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Best Local
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                                                                                                                                                                                                                                                         (complementarity determining regions) derived from a non-human antigen-
specific donor antibody, and an acceptor framework from a non-human
primate. The Abs are prepared by grafting CDRs from a non-human antigen-
specific donor antibody onto homologous Old World ape or monkey acceptor
frameworks. The Abs have reduced immunogenicity and are better tolerated
in humans (because of the close similarity between the human and primate
proteins), but retain the full antigen-binding affinity of the donor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypercalcaemia. The agent contained a portion of the antihuman parathyroid hormone-related protein monoclonal antibody (antihuman PTHFP MAb). The MAb was used as the active component in the agent. The agent further comprises a rodent/human chimeric MAb which has a rodent variable region and a human constant region and recognises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 26; Page 101; 123pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antibodies containing donor complementarity determining regions and non-human primate acceptor frameworks, having reduced immunogenicity humans -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-023265/02
N-PSDB; AAZ39338.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Complementarity determining region; antibody; primate; immunogenicity; Old World ape; Old World monkey; antigen-binding affinity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Partial peptide fragment of Cynomologous Vkappa cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY56670 standard;
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                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention provides an antibody (Ab) comprising donor CDRs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-NOV-1999
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1 FTLKISR 7
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                                                 Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
7; Conserv
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                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                    Ă,
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                                                   Score 33; DE
Pred. No. 3.8
0; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                            DB;
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                                                                                                     21;
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                                                      0,
                                                                                                     Length 93;
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RESULT
AAR72066
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                       AAR72066;
                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                              gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         L- and H-chain DNA was amplified by PCR from Graves' orbital
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 66; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Graves'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mclachlan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              variable region; autoimmunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OF7K.7 VK-2 L chain
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                                                                 AAR72066 standard;
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7; Conserv
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54.60
/label= CDR2
93..100
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(first entry)

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RESULT
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XX AAE1
XX AAE1
XX AAE1
XX AAE1
XX YURG
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Best Local S
Matches 7
           Unidentified
                             Tumour-associated antigen mucin-1; MUC-1; adenocarcinoma; variable light chain region; cancer; breast; ovary; lung; cytostatic; therapy.
                                                                                                                                                                                                                                                                                                                    L- and H-chain DNA was amplified by PCR from Graves' orbital tissue and clones encoding autoimmune-associated immunoglobulin fragments were obtained. 14/15 clones of L chain (kappa) regions showed homology to the putative VK germline gene KLO12 (given in AAQ89317) and the remaining clone, OFFK.7 (AAQ89326), to the VKO05
                                                                                             04-JAN-2002
                                                                                                                   AAE12711;
                                                                                                                                      AAE12711 standard;
                                                                                                                                                                                                                                                                                                                                                                                                        Graves' ophthalmopathy-associated monoclonal antibody - produced by molecular cloning of immunoglobulin genes by \operatorname{\mathsf{PCR}}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Graves ophthalmopathy associated immunoglobulin protein; orbital antigen; monoclonal antibody; light chain; L chain; variable region; autoimmunity.
                                                                      Antibody variable light chain region (VL)
                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                    Local Similarity nes 7; Conserv
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                                                                                                                                                                                                                 FTLKISR 7
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                                                                                                                                                                                                                                      Conservative
                                                                                             (first entry)
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93..100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers 23..38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= CDR3
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                                                                                                                                      peptide; 100
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Pred. No.
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RESULT 10
AAE06958
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated tumour-associated antigen mucin-1 (MUC-1)-specific binding member comprising an antigen binding domain (CC region having an antibody variable light (VL) or heavy (VH) region, CC or a complementarity determining region (CDR) of VL or VH. MUCI-specific binding member is useful for diagnosing cancer, preferably adenocarcinoma CC The binding of MUCI-specific binding member to MUCI is detected by a detection method selected from enzyme-linked immunosorbent assay, CC MUCI-specific binding member is useful for treating cancer, preferably cadenocarcinoma, in an individual, where the cancer is present in tissue of the breast, ovary, lung, or bladder of the individual. MUCI-specific binding member is useful for diagnosing and imaging MUCI-expressing CC content cells and tissues, for purifying or isolating non-glycosylated, conter cells and tissues, and for therapeutically or prophylactically containing molecules, and for therapeutically or prophylactically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                              multiple scierosis; atherogenesis; atherosclerosis; restenosis; asthma; anaphylaxis; malignancy; inflammation; stenosis; allograft rejection; fibrotic disease; angioplasty; acquired immune deficiency syndrome; AIDS; inflammatory glomerulopathy; vascular intervention; ID9 antibody; neointimal hyperplasia; VK; kappa light chain variable region.
                                                                                                                                                        Murine; humanised antibody; CC-chemokine receptor 2; CCR2; nephrotropic; neuroprotective; immunosuppressive; human immunodeficiency virus; HIV infection; cytostatic; vasotropic; leukocyte trafficking; allergy; inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shock;
                                                                                                                                                                                                                                                          Murine 1D9
                                                                                                                                                                                                                                                                                               16-OCT-2001
                                                                                                                                                                                                                                                                                                                                  AAE06958;
                                                                                                                                                                                                                                                                                                                                                                        AAE06958 standard; Protein; 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated tumor-associated antigen mucin-1-specific binding member for diagnosing and treating cancer, comprises mucin-1 binding domain or its portion for binding to an epitope of the protein core of mucin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-626437/72.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ng cancer. The present sequence is antibody variable light chain
(VL) from DP47 germ line.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 7; Conserv
                                                                                                                                                                                                                                                      antibody partial kappa light chain variable (VK) region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 35; 126pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; ilarity 100.0%; Conservative (
                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                           entry)
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Pred. No.
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RESULT J
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AC AAE
XX AAE
DT 16-C
DT 16-C
DT MOUS
XX MOUS
KW HIV
KW PIV
KW Infl
KW Anal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fragment, having binding specificity for CC-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The humanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating HV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as inflammatory disorder, autoimmune disorders such as resulting and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and in the manufacture of a medicament for treating CCR2-mediated disease. They are also useful for treating allergy, anaphylaxis, and lighted allergy, chronic and acute inflammation, histamine and IgE-mediated allergy chronic and acute inflammation, histamine and IgE-mediated allergy chronic and acute inflammation, histamine and IgE-mediated.
                      inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shock; multiple sclerosis; atherogenesis; atherosclerosis; restenosis; asthma; anaphylaxis; malignancy; inflammation; stenosis; allograft rejection;
                                                                               Mouse; humanised antibody; CC-chemokine receptor 2; CCR2; nephrot neuroprotective; immunosuppressive; human immunodeficiency virus; HIV infection; cytostatic; vasotropic; leukocyte trafficking; all
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity for CCR2, and a non-human antigen binding region and human immunoglobulin
                                                                                                                                                              Mouse germline kappa light chain variable (VK) region,
                                                                                                                                                                                                                                             AAE06959
                                                                                                                                                                                                                                                                                    AAE06959 standard; Protein;
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fibrotic
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nes 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                               ftlkisr 82
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        deficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 100;
                                                                                                                       CCR2; nephrotropic
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                                                                                     allergy;
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RESULT 1

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AAE06960 standard; Protein; 100

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BXSXFXBXB

16-0CT-2001 AAE06960;

(first entry)

germline

Mouse; Mouse

humanised antibody; CC-chemokine receptor 2; CCR2; nephrotropic;

kappa light chain variable (VK) region, 70/1.

Q

1 FTLKISR 7 ftlkisr

Query Match Best Local Similarity

100.0%;

Score 33; Pred. No.

DB 22; 0

Length 100;

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Gaps

0

Matches

Conservative

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Mismatches

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inflammatory disorder, autoimmune disorders such as rheumatoid arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and in the manufacture of a medicament for treating CCR-2 mediated disease. They are also useful for treating allergy, anaphylaxis, malignancy, chronic and acute inflammation, histamine and IgE-mediated allergic reaction, shock, stenosis, allograft rejection, fibrotic disease, asthma, inflammatory glomerulopathies, acquired immune deficiency syndrome (ALDS), restenosis associated with vascular intervention, including angioplasty and/or stent placement in a mammal. Humanised antibodies are also useful for inhibiting nerowing of the lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of a vessel in a mammal, preferably associated with vascular intervention.
                                                                                                                                                                                                                                                                                                                                (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The humanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating HIV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The patent discloses a humanised antibody or its antigen-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity for CCR2, and a non-human antigen binding region and human immunoglobulin
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Sequence
                                       The present sequence is (VK) region, 70/3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 147; 183pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Larosa
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                                                       germline kappa light
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RESULT
AAE06961
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AC AAE
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Best Local
                                                                                                                                                                                                                                                                                                                                                                           and in the manufacture of a medicament for treating CCR-2 mediated disease. They are also useful for treating allergy, anaphylaxis, malignancy, chronic and acute inflammation, histamine and IgE-mediated allergic reaction, shock, stenosis, allograft rejection, fibrotic disease, asthma, inflammatory glomerulopathies, acquired immune deficiency syndrome (AIDS), restenosis associated with vascular intervention, including angioplasty and/or stent placement in a mammal. Humanised antibodies are also useful for inhibiting narrowing of the lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of a vessel in a mammal, preferably associated with vascular intervention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The patent discloses a humanised antibody or its antigen-binding fragment, having binding specificity for Cc-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin. The humanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating HIV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as inflammatory disorder, autoimmune disorders such as rheumatoid arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and for inhibiting restenosis. They are useful in therapy or diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroprotective; immunosuppressive; human immunodeficiency virus; HIV infection; cytostatic; vasotropic; leukocyte trafficking; allergy; inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shoc multiple sclerosis; atherogenesis; atherosclerosis; restenosis; asthma anaphylaxis; malignancy; inflammation; stenosis; allograft rejection; fibrotic disease; angioplasty; acquired immune deficiency syndrome; AIDS; inflammatory glomerulopathy; vascular intervention; neointimal hyperplasia; VK; kappa light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity for CCR2, and a non-human antigen binding region and human immunoglobulin
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                                       AAE06961 standard; Protein; 100 AA.
                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                         Local Similarity res 7; Conserv
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|ftlkisr 82
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                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                               is mouse
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                                                                                                                                                                                                             0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                           germline kappa light chain
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                                                                                                                                                                                                                                                                                                                                                               variable
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CC humanised antibodies are useful for inhibiting the interaction of Cc a cell expressing CCR2. They are useful for inhibiting the interaction of Cc a cell expressing CCR2. They are useful for inhibiting or treating CC HIV infection. The proteins of the invention are useful for inhibiting CCR2. They are useful for inhibiting CCR2 mediated disorders such as CC inflammatory disorder, autoimmune disorders such as rheumatoid CC arthritis and multiple sclerosis, atherogenesis and atherosclerosis, CC and for inhibiting restenosis. They are useful in therapy or diagnosis, CC and in the manufacture of a medicament for treating CCR-2 mediated CCR2 mediated allergic reaction, shock, stenosis, allergy, anaphylaxis, CC malignarcy, chronic and acter inflammation, histamine and IgE-mediated allergic reaction, shock, stenosis, allograft rejection, CC fibrotic disease, asthma, inflammatory glomerulopathtes, acquired CC immune deficiency syndrome (AIDS), restenosis associated with vascular intervention, including angioplasty and/or stent placement in a mammal. CC Humanised antibodies are also useful for inhibiting narrowing of the CC usesel in a mammal, and inhibiting neointimal hyperplasia of CC The present sequence is mouse germline kappa light chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    multiple sclerosis; atherogenesis; atherosclerosis; restenosis; asthma; anaphylaxis; malignancy; inflammation; stenosis; allograft rejection; fibrotic disease; angioplasty; acquired immune deficiency syndrome; AIDS; inflammatory glomerulopathy; vascular intervention; neointimal hyperplasia; VK; kappa light chain variable region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         fragment, having binding specificity for CC-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity for CCR2, and a non-human antigen binding region and human immunoglobulin
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                                     The present sequence (VK) region, 70/2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MILL-) MILLENNIUM PHARM INC
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Query Match
Best Local Similarity
"hes 7; Conserva

Conservative

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Gaps

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100.0%;

Score 33; Pred. No. Mismatches

DB 22;

Length 100;

Sequence

100 AA;

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RESULT JABCO 6922 ID AAECO 5022 ID AAECO 5022 ID AAECO XX AAECO XX HOUSE KW HIV KW HIV KW AIDS KW HIV KW AIDS 
                                                                                                                                                                                                  The patent discloses a humanised antibody or its antigen-binding CC fragment, having binding specificity for CC-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin. The CC and at least a portion of an immunoglobulin of human origin. The CC humanised antibodies are useful for inhibiting the interaction of CC a cell expressing CCR2. They are useful for inhibiting or treating CCR inhibiting in the context of the invention are useful for inhibiting CCR inhibiting in the context of the invention are useful for thematoid CC arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and in the manufacture of a medicament for treating CCR-2 mediated CC disease. They are also useful for treating allergy, anaphylaxis, CC malignancy, chronic and acute inflammation, histamine and IgE-cc mediated allergic reaction, shock, stenosis, allograft rejection, CC fibrotic disease, asthma, inflammatory glomerulopathies, acquired CC intervention, including angioplasty and/or stent placement in a mammal. CC lumen of a vessel in a mammal, and inhibiting narrowing of the CC lumen of a vessel in a mammal, and inhibiting narrowing of the
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                                                                                                        lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of
a vessel in a mammal, preferably associated with vascular intervention.
The present sequence is mouse germline kappa light chain variable
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                                                                                                                             inflammatory disorder, autoimmune disorders such as rheumatoid arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and in the manufacture of a medicament for treating CCR-2 mediated disease. They are also useful for treating allergy, anaphylaxis, chronic and scutch inflammation, histamine and IgE-modification of the state of the s
                                                                                                                                                                                                                                                                                                   The patent discloses a humanised antibody or its antigen-binding fragment, having binding specificity for CC-hemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The humanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating HIV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shock multiple sclerosis; atherogenesis; atheroselerosis; restenosis; asthma; anaphylaxis; malignancy; inflammation; stenosis; allograft rejection; fibrotic disease; angioplasty; acquired immune deficiency syndrome; aIDS; inflammatory glomerulopathy; vascular intervention; neointimal hyperplasia; VK; kappa light chain variable region.
                                                mediated allergic reaction, shock, stenosis, allograft rejection, fibrotic disease, asthma, inflammatory glomerulopathies, acquired immune deficiency syndrome (AIDS), restenosis associated with vascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Humanized immunoglobulin 2-mediated disorder in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 149; 183pp; English
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HIV infection; c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse; humanised antibody; CC-chemokine receptor 2; CCR2; nephrotropic
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                         including angioplasty and/or stent placement
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cytostatic; vasotropic; leukocyte trafficking; allergy;
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CC lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of CC a vessel in a mammal, preferably associated with vascular intervention. CC The present sequence is mouse germline kappa light chain variable CC (VK) region, V-IC.

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SQ Sequence 100 AA;

Query Match
Best Local Similarity 100.0%; Score 33; DB 22; Length 100;
Best Local Similarity 100.0%; Pred. No. 4;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 FTLKISR 7

Qy 1 FTLKISR 7

Db 76 ftlkisr 82

Search completed: July 15, 2002, 12:57:58

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Minimum
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Listing first 45 summaries
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US-08-488-133B-151

US-08-881-037-37

US-08-881-13B-151

US-08-107-669D-15

US-08-177-689D-15

US-08-477-781B-15

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US-08-378-939-28

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ALIGNMENTS

RESULT 1 US-08-477-877B-33

Sequence 33, Application US/08477877B Patent No. 5730979

GENERAL INFORMATION:

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; TOPOLOGY: 1:; MOLECULE TYPE: US-08-477-877B-33
                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/407,009
EILING DATE: 29-MAR-1995
APPLICATION NUMBER: 08/119,032
FILING DATE: 09-SEP-193
APPLICATION NUMBER: 08/027,008
FILING DATE: 05-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: 01stein, Eliot M.
                                                                                                                                     TELEPHONE: 201-994-174
TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
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MEDIUM TYPE: 3.5 inc
COMPUTER: IBM PS/2
                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bazin, Herv
APPLICANT: Latinne, Dominique
TITLE OF INVENTION: LO-CD2a Antibody and Uses Thereof for Inhibiting T-Cell Act
NUMBER OF SEQUENCES: 96
                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: UZIP: 07068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 6 Beck
                                                         STRANDEDNESS:
                                                                                                                                                                                                                                      NAME: Olstein, Elliot
REGISTRATION NUMBER: 2
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                                                                                                LENGTH:
                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New Jersey
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                                                                                            32 amino acids
                                           linear
                    polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inch diskette
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Query Match
Best Local Similarity
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US-07-977-696C-71
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                                                                          Sequence 71,
Patent No. 5
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Best Local :
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                                                              GENERAL INFORMATION:
APPLICANT:
APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: peptide 7-977-696C-71
                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
LENGTH: 32 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 5.0
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/977,696C
FILING DATE: 11-16-92
                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: P6
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 748-6868
TELEPAX: (510) 748-6688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Amzel Ph.D., Viviana
REGISTRATION NUMBER: 30,93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Padlan Dr., Eduardo A. TITLE OF INVENTION: Analogue Pept TITLE OF INVENTION: for Carcinomation: and Therapeut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: do Couto, Fernando J.R.
APPLICANT: Ceriani Dr., Roberto L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Peterson Dr.,
                                                                                                                                                                 |||||||
15 FTLKISR 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 444 South Flower Street, Suite 2000 CITY: Los Angeles STATE: California COUNTRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: C
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                                                                                                                                                                                                1 FTLKISR 7
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5. 5792852
                                                                             l, Application US/08129930B
5804187
                                                                                                                                                                                                                                                                                                                                               amino acids
do Couto Dr., Fernando .
Ceriani Dr., Roberto L.
Peterson Dr., Jerry A.
Padlan Dr., Eduardo A.
                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                     100.0%;
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                                           Fernando J.R.
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Pred. No.
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INFORMATION FOR SEQ ID NO:
COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,281A
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                 APPLICANT: Bazin, Herv
APPLICANT: Latinne, Dominique
TITLE OF INVENTION: LO-CD2a Antibody and Uses Thereof for Inhibiting T-Cell Act
NUMBER OF SEQUENCES: 96
CORRESPONDENCE ADDRESS:
                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 521-133
TELEPAX: (510) 521-3541
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CORRESPONDENCE ADDRESS:
ADDRESSEE: V. AMZEL & ASSOC.
STREET: 2055 No. 5804187th Broadway, Suite 201
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                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                              STATE:
                                                                                                                                                                                                         CITY: Roseland
                                                                                                                                                                                                                               STREET:
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ADDRESSEE: Cecchi, Stewart & Olstein
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OPERATING SYSTEM:
SOFTWARE: PatentI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94596
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SYSTEM: PC-DOS/MS-DOS 5.
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Pred. No. 0.72;
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Best Local Similarity
Watches 7; Conserve
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APPLICATION NUMBER: 08/407,009
FILING DATE: 29-MAR-1995
APPLICATION NUMBER: 08/219,032
FILING DATE: 09-SEP-1993
APPLICATION NUMBER: 08/027,008
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: 08/027,008
FILING DATE: 05-MAR-1993
APPLICATION TIPORMATION:
NAME: 01stein, Elliot M.
REGISTRATION NUMBER: 24,025
REFERNATION NUMBER: 61750-142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Postema, Christina E. APPLICANT: White-Scharf, Mary TITLE OF INVENTION: CD-CD2a Antil TITLE OF INVENTION: Thereof for TITLE OF INVENTION: T-Cell Active TITLE OF INVENTION: Proliferation
                                                                                    APPLICATION NUMBER: US/08/477,989B FILING DATE: 07-JUN-1995 CLASSIFICATION: 424 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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            APPLICATION NUMBER: 08/11 FILING DATE: 09-SEP-1993
                                                                                                                                                                          OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
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STATE: New Jer:
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 FILING DATE: 09-SE APPLICATION NUMBER:
                                                   APPLICATION NUMBER: 08/407,009 FILING DATE: 29-MAR-1995
                                                                                                                                                                                                               COMPUTER:
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o. 5951983
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                                  08/119,032
08/027,008
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US-08-976-288A-71
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INFORMATION FOR SEQ ID NO: 33
SEQUENCE CHARACTERISTICS:
FENGTH: 32 amino acids
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APPLICANT: do Coute
APPLICANT: Cerianti
APPLICANT: Peterso:
APPLICANT: Padlan i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
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                                                                TELEX: n.a.
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: FITLE OF INVENTION: ITTLE 
                                                                                                                                                                 NAME: Viviana Amzel Ph.D.
REGISTRATION NUMBER: 30,930
REFERENCE/DOCKET NUMBER: P6639938
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 622-7700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/129,930
FILING DATE: September 30, 1993
PAIOR APPLICATION DATA:
APPLICATION NUMBER: 07/977,696
FILING DATE: NO. 6315997ember 16, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Viviana Amzel Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 5.0
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                       SEQUENCE CHARACTERISTICS:
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TOPOLOGY: linear
MOLECULE TYPE: poly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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NAME: Olstein, Elliot M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                        TELEFAX:
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15 FTLKISR 21
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REFERENCE/DOCKET NUMBER: 61
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California
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32 amino acids
                                                                                                                                        (213) 489-4210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          do Couto Dr., Fernando Ceriani Dr., Roberto L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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No. 6315997ember 21, 1997
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                                                                       71:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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Pred. No.
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; MOLECULE TYPE: US-08-976-288A-71
RESULT 8:
US-08-881-037-37
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Best Local Similarity
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Best Local Similarity 100.0%;
Matches 7; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/497,312
FILING DATE: 30-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
REGISTRATION NUMBER: 30.549
REFERENCE/DOCKET NUMBER: 26:9US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 801/532-1922
                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 801/531-9168
TELEX: 388961 1PM04UT
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 81 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
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TITLE OF INVENTION: Immunoglobulins with reduced immunogenicity of murine
TITLE OF INVENTION: Intrody variable domains, compositions containing them.
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: CENTRO DE INMUNOLOGIA MOLECULAR
STREET: 215 Y 15, ATABEY PLAYA
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MEDIUM TYPE: Floppy disk
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HYPOTHETICAL:
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 11600
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                                                                           53 FTLKISR 59
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5712120
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                                                                                                                                                                                                                                                                            unknown
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Pred. No. 0.72;
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Pred. No. 1
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US-08-488-113B-151
; Sequence 151, Application US/08488113B
; Patent No. 5744580
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Best Local Similarity
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                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Better, Marc D.
APPLICANT: Carroll, Stephen F.
APPLICANT: Studnika, Gary M.
TITLE OF INVENTION: Immunotoxins
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 169
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 37, Application US/08881037
Patent No. 6080588
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 23-JUN-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 18-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: KONSK1, ANTOINETTE F.
REGISTRATION NUMBER: 34,202
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELECHIONE: (650) 813-5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/881,037
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESSE: Morrison & Foerster
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Glick, Gary D.
APPLICANT: Swanson, Patrick C.
TITLE OF INVENTION: DNA BINDING ANTIBODIES
COUNTRY: USA
ZIP: 60661
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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CITY: Palo Alto
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STRANDEDNESS: sir
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                                                                              STATE:
                                                                                                                 STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94304-1018
                                                                                                                                 ADDRESSEE:
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                                                                            Chicago
: Illinois
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ilarity 100.0%;
Conservative (
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   Floppy disk
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CURRENT APPLICATION DATA:

SOFTWARE:

PatentIn Release #1.0, Version #1.25

APPLICATION NUMBER:

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

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RESULT 10
US-08-477-484B-151
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Best Local Similarity
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US-08-488-113B-151
                                                                                                                                                                                                                                                                                                      Sequence 151, Application US/08477484B Patent No. 5756699
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                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                               TITLE OF INVENTION: Immunoto:
                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                          APPLICANT: Studnika, Gary M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/988,430
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/901,707
FILING DATE: 19-JUN-1992
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 12-MAY-:

PRIOR APPLICATION DATA:
                                                                                                                                           CITY: Chicago
                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                                                                                                       72 FTLKISR 78
APPLICATION NUMBER:
                                                                                                             COUNTRY:
                                                                                                                                                                      ADDRESSEE:
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TELEFAX: 650 388-1248
TELEX: 650 388-1248
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                                                                                                                                                                                                                                                                                                                                                                                                                       1 FTLKISR 7
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                                                                                                   60661
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                                                                                                                            Illinois
                                                                                                                                                    E: McAndrews, Held & Malloy, Ltd.
                                                                                                               USA
                                                                                                                                                                                                                                                        Better, Marc D.
Carroll, Stephen F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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07-JUN-1995
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12-MAY-1993
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US/08/477,484B
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 650 388-1248 INFORMATION FOR SEQ ID NO:
                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
              CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
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REFERENCE/DOCKET NUMBER: 11022US07/200-70.P3.C2A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-9155
TELEZX: 312/707-9155
TELEX: 650 380-111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: MCNicholas, Janet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/988,430
FILING DATE: 09-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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                                                                                                                APPLICATION NUMBER: FILING DATE: 13-AUG
                                                                                                                                                                                                                                                                                               CITY: Washington
                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            72 FTLKISR 78
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                                                                                                                                                                                                                                                                                                                 E: Sterne, Kessler,
1100 New York Ave.,
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                                                                                                                13-AUG-1993
JMBER: US 07/808,464
13-DEC-1991
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12-MAY-1993
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                                                                                                                                                                                                                                                                                                                                                                     Modified Antibody Variable Domains (as amended): 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                us/08/107,669p
                                                                                                                                                                                                                                                                                                                                                                                                   Gary M.
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                                                                                                                                                                                                                                                                                                               Goldstein and Fox P.L.L.C N.W., Suite 600
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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-107-669D-15
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US-08-472-788A-15
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SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino ****
                                                                                                                                                                                                                           APPLICATION NUMBER: 08/082,842
EILING DATE: 23-JUN-1993
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION UMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA: US 07/808,464
EILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele 32,651
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                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 202/371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,788A
FILING DATE: 07-JUN-1995
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                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acid
                                                                                                                                                          REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0610.1000003
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2600
                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                  MOLECULE TYPE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                   TOPOLOGY:
                                                                                                                                           TELEFAX:
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                                                                   108 amino acids
                                                                                                                                           202/371-2540
                                   linear
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RESULT 13
US-08-477-531B-15
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문
                                                                                                                                                ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-477-531B-15
                                                  Query Match
Best Local Similarity
"hehes 7; Conservations
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Best Local S
Matches 7
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                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/107,669
FILING DATE: 13-AUG-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
APPLICATION NUMBER: US 07/808,464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                        TELEFAX: 202/371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0610.1000004/MAC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2600
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acid
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72 FTLKISR 78
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                                 1 FTLKISR 7
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o. 5821123
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1100 New York Ave.,
                                                                                                                                                                                                                      108 amino acids
                                                                  Conservative
                                                                                                                                                                                                                                                                          202/371-2540
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                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Release #1.0, Version #1.25
                                                                   0
                                                                  Score 33; DB Pred. No. 2.3; Mismatches
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                                                                                                    DB 2;
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RESULT 14
US-08-646-360-151
Sequence 151, Application US/08646360
; Patent No. 5837491

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RESULT 15
US-08-082-842A-15
; Sequence 15, Application US/08082842A; Patent No. 5869619
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                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 7
         GENERAL INFORMATION:
APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified Antibody Variable Domains
NUMBER OF SEQUENCES: 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US94/05348
FILING DATE: 12-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/988,430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 650 388-1248 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENERAL INFORMATION:
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LENGTH: 108 amino acids
TYPE: amino acid
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: PCT/US94/05348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 200-70.P4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-8889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/901,707
FILING DATE: 19-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 173
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/787,567
EILING DATE: 04-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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CITY: Chicago
STATE: Illinoi
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                                                                                                                                                                                                                                                                            Local Similarity les 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 312/707-888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: McNicholas, Janet M. REGISTRATION NUMBER: 32,918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/01
FILING DATE: 13-MAY-1996
                                                                                                                                                                                                                           1 FTLKISR 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3: McAndrews, Held & Malloy, Ltd.
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Carroll, Stephen F.
Studnika, Gary M.
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Search completed: July 15, Job time: 390 sec
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                                                                                                                                                Query Match
Best Local Similarity
Matches 7; Conserv
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 06
TELECOMBUNICATION INFORMATION:
TELEPHONE: 202/371-2600
TELEFAX: 202/371-2540
                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                           TYPE:
                                                                                   72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 23-JU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Sterne, Kessler, Goldstein and Fox P.L.L.C. STREET: 1100 New York Ave., N.W., Suite 600 CITY: Washington
                                                                                                        1 FTLKISR 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER:
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                                                                                   FTLKISR 78
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Minimum
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Perfect score:
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Match Length
   PIR_71:*
1: pir1:*
2: pir2:*
3: pir3:*
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Gapop 10.0 , Gapext
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Listing first
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   GenCore version (c) 1993 - 2000
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kappa chain
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kappa	-		light	kappa	-	kappa	light	kappa	kappa	heavy	kappa	kappa	kappa		

ALIGNMENTS

Ig kappa chain V-III region (CP1) - hum C;Species: Homo sapiens (man) C;Date: 23-Oct-1992 #sequence_revision C;Accession: JT0521

human (fragment)

23-Oct-1992 #text_change 09-May-1997

patients with x-linked agammagl

ij

chain

rearrangeme

RESULT

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R; Anker, R; Conley, M.E.; Pollok, B.A.
J. Exp. Med. 169, 2109-2119, 1989
A; Title: Clonal diversity in the B cell repertoire of patients wi
A; Reference number: JT0511; MUID:89279157
A; Accession: JT0521
A; Mclecule type: mRNA
A; Mclecule type: mRNA
A; Residues: 1-54 <ANN'>
A; Note: the sequence shown here is one of eight productive V-D-J
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 1-49/Domain: V region <VRE>
F; 50-54/Domain: J region <JRE>
                                                                                                                                                                                                                                                                                                                                     Ig kappa chain V region - human (fragment)
(;Species: Homo sapiens (man)
(;Species: Homo sapiens (man)
(;Date: 20-Feb-1995 #sequence_revision 25-Oct-1996 #text_change 23-Jul-1999
(;Accession: $34082; $21526
R;Wagner, S.D.; Luzzatto, L.
R;Wagner, S.D.; Luzzatto, L
                                                                                                                                                                                                           A;Cross-references: EMBL:X66042; NID:g33318; PIDN:CAA46841.1; PID:g33319 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
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                                             Score 43; DB 2;
Pred. No. 0.058;
    Mismatches
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Ig kappa chain - mouse (fragment)
C;Species; Mus musculus (house mouse)
C;Date: 19-Feb-1994 *sequence_revision 10
C;Accession: S24215
R;Shimizu, T.; Iwasarn "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig kappa chain V region - human
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Ja
C;Accession: S34102
R;Wagner, S.D.; Luzzatto, L.
Eur. J. Immunol. 23, 391-397, 1993
Eur. J. Immunol. 23, 391-397, 1993
A;Title: V-kappa gene segments rearranged in
A;Reference number: S34076; MUID:93170387
A;Accession: S34102
                                                                                                                                                                                                                                                                                                                                                                                                                                                    R; Snimizu, T.; Iwasato, T.; Yamagishi, H.
J. Exp. Med. 173, 1065-1072, 1991
A; Title: Deletions of immunoglobulin C(kappa)
A; Reference number: S24214; MUID:91217618
A; Accession: S24215
 Ig light chain V region (clone 165.49) - mouse (fragment) C;Speciles: Mus musculus (house mouse) C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change C;Accession: PH1048
R;Tillman, D.M.; Jou, N.T.; Hill, R.J.; Marion, T.N.
J. Exp. Med. 176, 761-779, 1992
A;Title: Both IgM and IgG anti-DNA antibodies are the products of Reference number: PH0971; MUID:92381444
A;Accession: PH1048
A;Status: nucleic acid sequence not shown
                                                                                                                                                                                                                                                                                                                                                              A:Cross-references: EMBL:x58202; NID:g53718; PIDN:CAA41178.1; PID:g930195 C:Superfamily: immunoglobulin V region; immunoglobulin homology C:Keywords: heterotetramer; immunoglobulin
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-79 <SHI>
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A; Residues: 1-78 <WAG>
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A;Cross-references: EMBL:X54832
C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
E;7-81/Domain: immunoglobulin homology <IMM>
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A:Residues: 1-86 <WAGS
A:Cross-references: EMBL:X67185
C:Superfamily: immunoglobulin V region; immunoglobulin homology
C:Keywords: heterotetramer; immunoglobulin
E:7-81/Domain: immunoglobulin homology <IMM>
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A.Molecule type: mRNA
A.Molecule type: mRNA
A.Residues: 1-86 <BLA>
A.Cross references: EMBL:X54824; NID:g33653; PIDN:CAA38593.1; PID:g33654
A.Cross references: EMBL:X54824; NID:g33653; PIDN:CAA38593.1; PID:g33654
R:Wagner, S.D.; Luzzatto, L.
Eur. J. Immunol. 23, 391-397, 1993
A.Title: V-kappa gene segments rearranged in chronic lymphocytic leukemia are distrib
A.Reference number: S34076; MUID:93170387
A.Reference number: S34011
A.Status: preliminary
A.Status: preliminary
                                                                                     A;Status: preliminary; translation
A;Molecule type: mRNA
A;Residues: 1-86 <BLA>
                                                                                                                                                R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L. Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III
A;Reference number: S16823; MUID:91243737
A;Accession: S16834
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(Speckes: Homo sapiens (man)
(C;Date: 19-reb-1994 #sequence_revision 10-Nov-1995
(C;Accession: S16834
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S16834
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Thehes 9; Conserv
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C;Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S16826; S34101
R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal r)
A;Reference number: S16823; MUID:91243737
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Ig Kappa Chain V region - human
C:Species: Homo sapiens (man)
C;Date: 13-Jan-1995 *sequence_revision 13-Jan-1995 *text_change 21-Jan-2000
C;Accession: S34086
R;Wagner, S.D.; Luzzatto, L.
Eur. J. Immunol. 23, 391-397, 1993
A;Mitle: V-kappa gene segments rearranged in chronic lymphocytic leukemia are d
A;Reference number: S34076; MUID:93170387
A;Accession: S34086
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-66 <WAG>
A;Cross-references: EMBL:X67169
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;9-83/Domain: immunoglobulin homology <IMM>
RESULT 10
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If kappa chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 *sequence_revision 10-Nov-1995 *text_change
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C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;7-81/Domain: immunoglobulin homology <IMM>
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R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable
A;Reference number: S16823; MUID:91243737
A;Accession: S16836
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A; Residues: 1-86 <BLA>
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C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 *sequence_revision 10-Nov-1995 *text_change 21-Jan-2000
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Pred. No. 0.071;
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 *sequence_revision 10-Nov-1995 *text_change 21-Jan-2000
C;Accession: S16833; S16838
R;Blaison, G; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid A;Reference number: S16823; MUID:91243737
A;Accession: S16833
A;Accession: S16833
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S16833
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(;Species: Homo sapiens (man)
(;Species: Homo sapiens (man)
(;Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
(;Accession: S1637
R;Blaison, G; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Inmunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid A;Reference number: S1682; MUID:91243737
A;Reference number: S1682; MUID:91243737
A;Accession: S1637
A;Accession: S1637
A;Molecule type: mRNA
A;Residues: 1-86 <BLA>
A;Residues: 1-86 <BLA
A;Residues: 1
                                                                       A:Molecule type: mRNA
A:Residues: 1-86 <BLA>
A:Cross-references: EMBL:X54831
A:Experimental source: clone bkv17 A:Accession: S16838
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;7-81/Domain: immunoglobulin homology <IMM>
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C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid A;Reference number: S16823; MUID:91243737
A;Accession: S16840
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A; Residues: 1-86 <BLA>
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Best Local S
Matches 9
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Pred. No. 0.071;
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Pred. No.
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A; Molecule type: A; Residues: 1-86

A; Status: translation not shown

mRNA

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RESULT 14

Ig kappa chain V region - human (fragment)
C. Species: Homo sapiens (man)
C. Species: Homo sapiens (man)
C. Species: Homo sapiens (man)
C. C. Accession: S16824
R. Blaison, G.; Kuntt, J.L.; Pasquali, J.L.
Bur. J. Immunol. 21, 1221-1227, 1991
A.; Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid fac A.; Reference number: S16823; MUID:91243737
A.; Residues: preliminary; translation not shown
A.; Koccession: S16824
A.; Residues: 1-86 SBLA>
A.; Cross -references: EMBL:X54822
C.; Superfamily: immunoglobulin V region; immunoglobulin homology
C.; Keywords: heterotetramer; immunoglobulin homology
C.; Keywords: immunoglobulin homology <IMM>
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Bur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of particle. Molecular analysis of V(kappa)III variable regions of particle. Since the strength of the strength o
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A;Experimental source: clone slkvl4
C;Superfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;7-81/Domain: immunoglobulin homology <IMM>
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C;Speciles: Homo sapiens (man)
C;Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S16830
R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
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S16830
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Matches 9; Conserv
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TDFTLTISS 68
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9; Conserv
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ilarity 100.0%;
Conservative 0
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Pred. No. 0.0
0; Mismatches
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Pred. No.
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Pred. No. 0.071;
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Ig kappa chain V region - human (fragment)
C; Species: Homo sapiens (man)
C; Species: 19-Feb-1994 *sequence_revision 10-Nov-1995 *text_change 21-Jan-2000
C; Accession: S16829
R; Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rh
A; Reference number: S16829
A; Accession: S16829
A; Status: preliminary; translation not shown
A; Status: preliminary; translation not shown
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C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin E;7-81/Domain: immunoglobulin homology <IMM>
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S16829
Search completed: July 15, 2002, 13:01:02 Job time: 468 sec
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A; Residues: 1-86 <BLA>
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Best Local Similarity
Matches 9; Conser
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ALIGNMENTS

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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation update)
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InterPro; IPR003596; Ig_v.
Pfam; PP00047; ig; 1.
SMART; SM00406; IGv; 1.
Immunoglobulin V region; Glycoprotein.
CARBOHYD 28 28 N-LINKED
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. NCBI_TaxID=9606; [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=75075135; PubMed=4216454;
Milstein C.P., Deverson E.V.;
"Primary structure of kappa light chain from
protein.";
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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; P80362; 1WTL.
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SCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE INV (1,2)
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107 AA;
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I region CAR.
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RESULT 3

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p1-1986 (Rel. 01, Created)
p1-JUL-1986 (Rel. 01, Last sequence update)
p15-JUL-1999 (Rel. 38, Last annotation update)
p15-JUL-1999 (Rel. 38, Last annotation update)
p16 kappa chain V-V region UPC 61.
p17 kappa chain V-V region UPC 61.
p18 musculus (Mouse).
p18 musculus (Mouse).
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Hoppe-Seyler's 2. Physiol. Chem. 351:1291-1295(1970).
-!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE
-!- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
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"The primary structure of a monoclonal kappa-type immunoglobulin L-
                                                                                                                                        DOMAIN
DOMAIN
                                                                                                                                                                                                HSSP; P80362; 1WTL.
Interpro; IPR003006; Ig_MHC.
Interpro; IPR003596; Ig_v.
pfam; PF00047; 1g; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                          Vrana M., Rudikoff S., Potter M.;
"The structural basis of a hapten-inhibitable kappa-chain idiotype.";
J. Immunol. 122:1905-1910(1979).
-i- MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM MYELOMA PROTEINS THAT
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HSSP; P80362; 1WTL.
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SMART; SM00406; IGv;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                       DOMAIN
                                                                                                                                                                                 Immunoglobulin
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24 34 COMPLEMENTARITY-DI
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
BY SIMILARITY.
                                       FRAMEWORK-1.

COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.

COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.

COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
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RESULT 4
KV5R_MOUSE
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P01651;
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21-JUL-1986 (Rel. (
15-JUL-1999 (Rel. 3
                                            KV5S_MOUSE STAND
P01652;
21-JUL-1986 (Rel. 01
21-JUL-1986 (Rel. 01
15-JUL-1999 (Rel. 38
15-JUL-1999 (Rel. 38
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SEQUENCE
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DISULFID
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J. Immunol. 122:1905-1910(1979).
-i- MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM MYELOMA PROTEINS THAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
Mus musculus (Mouse).
Eukaryota, Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
[1]
                                                                                                                                                                                                                                                                                                                     Pfam; PF00047; ig; SMART; SM00406; IG
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InterPro; IPR003596;
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.

COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
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Pred. No. 0.015;
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                  Craniata; Veri
Sciurognathi;
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                     Muridae;
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Johnson N., Slankard J., Paul L., Hood L.;
"The complete V domain amino acid sequences of
        DOMAIN
DISULFID
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DOMAIN
                                                                                                                                                                  MEDLINE-82099361; PubMed-6798111;
Johnson N., Slankard J., Paul L.,
"The complete V domain amino acid
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Mammalia; Eutheria;
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InterPro; IPR003596;
Pfam; PF00047; ig; 1
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                                                                                                                                                          binding proteins."
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                                                                                  SMART;
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                                                                                                                                 Immunol. 128:302-307(1982).
MISCELLANEOUS: THIS CHAIN WAS
BIND BETA(2-1)-FRUCTOFURANOSYL
                                                                                  SM00406;
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9; Conserv
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IPR003596;
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Rodentia;
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              FRAMEWORK - 1.

COMPLEMENTARITY - DETERMINING - 1.
FRAMEWORK - 2.
COMPLEMENTARITY - DETERMINING - 2.
FRAMEWORK - 3.

COMPLEMENTARITY - DETERMINING - 3.
FRAMEWORK - 4.
FRAMEWORK - 4.
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Pred. No.
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COMPLEMENTARITY - DETERMINING - 2.
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COMPLEMENTARITY-DETERMINING-1.
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Sciurognathi; Muridae;
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RESULT 8
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KV1T_H
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                                 KVIA_MOUSE STANDARD; PF P01632; Po1632; Po1632; Po1632; Po17986 (Rel. 01, Last seque 15-JUL-1986 (Rel. 01, Last annot 15-JUL-1999 (Rel. 38, Last annot 15 Rappa chain V-I region S107A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EULITZ M., Linke R.P.;

"Primary structure of the variable part of an amyloidogenic Bence-Jones Protein (Mev). An unusual insertion in the third hypervariable region of a human kappa-immunoglobulin light chain.";

HOPPE-Seyler's Z. Physiol. Chem. 363:1347-1358(1982).

-I- MISCELLANEOUS: ANOTHER FORM THAT LACKED PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P01612;
21-JUL-1986 (Rel.
21-JUL-1986 (Rel.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUMAN
 Eukaryota;
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                       Mus musculus
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nilarity 100.
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Pred. No. 0.015;
; Mismatches
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Best Local S
Matches 9
21-JUL-1986 (Rel. 01, Created)
21-JUL-1996 (Rel. 34, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 kappa chain v-1V region Len.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Ver
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                                                                                                                                                                                                                     Schneider M., Hilschmann N.;
"The primary structure of a monoclonic immunoglobulin-L-chain of subgroup IV of the kappa type (Bence-Jones protein Len).";
Hoppe-Seyler's Z. Physiol. Chem. 356:507-557(1975).
                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
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MEDLINE-81241357; PubMed-6788890;
Kwan S.-P., Rudikoff S., Seldman J.G.,
"Nucleic acid and protein sequences of
                                                                                                                                                                                                                                                                                                                                 MEDLINE-76004342; PubMed-50995;
                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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                                                                                                                                                                     REVISION TO 9.
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NCBI_TaxID~10090;
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
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                                                                                                                                  DOMAIN
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P01789;

.CPro; IPR003
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SMARY; SM00406; IGv; 1.

IMMUNoglobulin V region; Be.
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P13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
77 Fanna chain V-III region VG precursor ()
                                                                                                                                                                                                                                                                                                                                                         modified and this statement is not removed. entitles requires a license agreement (See or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                     the European Bioinformatics Institute. There are no rest use by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     within the VK locus.";
Nucleic Acids Res. 12:9229-9236(1984).
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InterPro; IPR003596; Ig_v.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                   IG KAPPA CHAIN V-III REGION VG
FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3
BY SIMILARITY.
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COMPLEMENTARITY-DETERMINING-1.
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Pred. No. 0.016;
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RESULT 11

KV3J_HUMAN

ID V3J_H

AC P04434

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KV3J_HUMAN STANDARD; PRT; 116 AA p04434; 13-AUG-1987 (Rel. 05, Created) 13-AUG-1987 (Rel. 05, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) 15 kappa chain V-III region VH precursor (F
                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR003006; Ig_MH
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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SIGNAL
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pech M., Zachau H.G.;
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9; Conserv
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Acids Res. 12:9229-9236(1984).
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      STANDARD;
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Pred. No. 0.0
Mismatches
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FRAMENCK-1.
COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
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BY SIMILARITY.
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Pred. No.
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 RESULT 13
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ID KV40_H
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A21056; A21056.
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Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
Fmmunoglobulin V region;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.

MEDLINE=81098966; PubMed=6779204;

Bentley D.L., Rabbitts T.H.;

"Human immunoglobulin variable region

kappa genes and a pseudogene.";

Nature 288:730-733(1980).
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain V-I region HK101 precursor (Fragment).
Homo sapiens (Human)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae,
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llarity 100.0%;
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BY SIMILARITY.
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Pred. No. 0.016;
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RESULT 14

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InterPro; IPR003596; IQ
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SMART; SM00406; ICV; 1
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MEDILINE-85014148; PubMed-6091049;

Klobeck H.G., Combriato G., Zachau H.G.;

Klobeck H.G., Combriato G., Zachau H.G.;

Klomunoglobulin genes of the kappa light cl

lymphoid cell lines are closely related.";

Nucleic Acids Res. 12:6995-7006(1984).
                                                                                                          Homo sapiens (Human).
Eukaryota; Metazoa; (
Mammalia; Eutheria; F
                                                                                                                                          13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain v-l region Walker precursor.
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entities requires a license agreement (See http://www.isb-sib.ch/announce/
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Homo sapiens
                                                                                               NCBI_TaxID=9606;
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9; Conserv
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Primates;
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COMPLEMENTARITY-DETERMINING-3
BY SIMILARITY.
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COMPLEMENTARITY-DETERMINING-2.
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Pred. No.
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FRAMEWORK-1.
                                                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Klobeck H.G., Bornka
Zachau H.G.
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01-JAN-1988 (Rel. 01-JAN-1988 (Rel. 01-JAN-1988 (Rel. 15-JUL-1999 (Rel. 15-J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Cranlata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                          HSSP;
                             [nterPro;
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V-IV region JI precursor.
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nkamm G.W., Comb
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COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
BY SIMILARITY.
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COMPLEMENTARITY-DETERMINING-1.
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DR Pfam; PF00047; ig; 1.

DR SMART; SM00406; IGV; 1

KW Immunoglobulin V region; S1>
FT CHAIN 21 133 IG N.
FT DOMAIN 44 60 COMPLEMEN N.

DOMAIN 61 75 FRAMEWORK 2.

POMAIN 61 75 COMPLEMENTARIT:

115 122 COMPLEMENTARIT:

123 132 FRAMEWORK 4.

43 114 BY SIMILARITY.

131 33 SCOTE 43; DB 1; Length Pred. No. 0.018;

Mismatches 0;
Search completed: July 15, 2002, 13:23:33
Job time: 1449 sec
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95 TDFTLTISS 103
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
BY SIMILARITY.
                                                                                                                   ; Score 43; DB 1; Length 133; ; Pred. No. 0.018; 0; Mismatches 0; Indels
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Listing first 45 summaries
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Q9u181 homo sapien
Q9u183 homo sapien
Q9u185 homo sapien
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Q94mb9 bacteriopha		Q94mc2 bacteriopha		Q94mg5 bacteriopha	_	S		Q26627 strongyloce	Q9h7w7 homo sapien	homo	bruc	homo sap	homo	Q9kte4 vibrio choi	09v2t0 pyrococcus	Q925s9 mus musculu		P74440 synechocyst	Q9a579 caulobacter	Ç,	Q9at27 phytophthor	4	S	000814 tritrichomo	Q99m37 mus musculu	mus	nus	Q9u180 homo sapien

ALIGNMENTS

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976 SAP

976 SAP

976 SAP

110 QS

121 QS

121 QS

122 QS

123 QS

124 QS

125 QS

125 QS

126 QS

127 QS

127
RESULT
Q9ERZ9
ID Q9
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Best Local S
Matches
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Q96SA9;
Q1-DEC-2001 (TrembLrel. 19, Created)
Q1-DEC-2001 (TrembLrel. 19, Last sequence update)
Q1-DEC-2001 (TrembLrel. 19, Last annotation update)
ANTI-STREPTOCOCCAL/ANTI-MYOSIN IMMUNOGLOBULIN KAPPA LIGHT CHAIN
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NON_TER
SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Adderson E.E., Shikhman A.R., Ward K.E., Cunningham M.W.; "Molecular analysis of polyreactive monoclonal antibodies from rheumatic carditis: human anti-N-acetylglucosamine/anti-myosin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. MEDLINE-98375893; PubMed-9712075;
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       Q9ERZ9
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J. Immunol. 161:2020-2031(1998).
EMBL; U96396; AAB68785.1; -
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9; Conservative
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107
           PRELIMINARY;
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11520 MW;
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Pred. No.
           PRT;
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Best Local S
Matches 9
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InterPro: IPR003596; Ig_V.
Pfam; PP00047; Ig; 1.
SMART; SM00409; IG; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IG; 1.
NON_TER 107 107
SEQUENCE 107 AA; 11784 MW;
                                                                                                                                                                                                                                                                                Q9UL77
Q9UL77;
01-MAY-2000
                                               Clin. Immunol. Immunopathol. EMBL; AF035037; AAD56273.1; - HSSP; P01607; IREI.
                                                                                                                                                                                                                                       01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                         Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Cloning genes of J. Cell. [2]
 InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                         SEQUENCE FROM N.A.
WEDLINE-98277139; PubMed-9614934;
WU X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.

Chen P., Deng J.B., Wang Z.L.,

Submitted (MAY-2000) to the EM

EMBL; AF262753; AAG23804.1; -.

HSSP; P80362; 1WTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.

Chen P., Deng J.B., Wang Z.L., Han H., Yao L.B.
"Construction and sequencing of the single-chair
human TNF-alpha specific monocional antibody.";
Ti. 4 Chun i Ta Hsueh Hsueh Pao 19:373-376(1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. Chen P., Deng J.B., Wang "Cloning and sequencing genes of an anti-hTNF-a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O9ERZ9;
O1-MAR-2001 (TrEMBLrel. 16, Created)
O1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI HUMAN TNF-ALPHA LIGHT CHAIN VARIABLE REGION (FI
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
NCBI_TaxID-10090;
                                                                                           fetus.
                                                                                                                                                                              NCBI_TaxID=9606;
                                                                                                                                                                                                                                 (FRAGMENT)
                                                                                                                 Young D.C
                                                                                                 Myosin-reactive autoantibodies
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Similarity 100.0%;
9; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IPR003599;
                                                                                                                                                                                                                                                                                                        PRELIMINARY;
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EMBL/GenBank/DDBJ databases.
                                                                          87:184-192(1998)
                                                                                                                                                                                       Craniata; Ve
Catarrhini;
                                                                                                                                                                                                                                                                                                        PRT;
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Pred. No. 0.16;
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Sciurognathi; Muridae;
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he single-chain
al antibody.";
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i; Hominidae;
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antibody
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Q9UL70;
01-MAY-2000 (Tremblrel. 13, C;
01-MAY-2000 (Tremblrel. 13, L;
01-DEC-2001 (Tremblrel. 19, L;
MYOSIN-REACTIVE IMMUNOGLOBULI)
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Q9UL81;
Q1-MAY-2000
Q1-MAY-2000
Q1-DEC-2001
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InterPro; IPR003596; 1
Pfam; PF00047; 1g; 1,
SMART; SM00406; IGV; 1
   Wu X.
               SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
Wu X., Liu B., Van der Merwe P.L.
                                                                                                AYOsı...
(FRAGMENT).
Homo sapiens (Human).
Homo sapiens (Human).
'horta; Metazoa; Chordata; 'horta; Primates;
                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                       01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clin. Immunol. Immunopathol. EMBL; AF035044; AAD56280.1; HSSP; P01607; 1REI.
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69 TDFTLTISS
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9; Conserv
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Milarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBLrel. 13, Last sequence update)
EMBLrel. 19, Last annotation update)
IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PubMed=9614934;
n der Merwe P.L.,
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108
11738 MW;
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                       P.L.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 43;
Pred. No.
                                                                                                                Craniata; Vo
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87:184-192(1998)
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Pred. No. 0.1
); Mismatches
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Catarrhini; Hominidae
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                                                                                                                                                                                         CHAIN VARIABLE
                                                                                                                                                                                                                                                                                          107
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                                                                                                                                Vertebrata; Euteleostomi;
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                   N.N.,
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                                                                                                                Hominidae;
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RESULT
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AC Q9
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Best Local S
Matches 8
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Best Local S
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Q9UL79 PRELIMINARY; PRT; 108 AA.
Q9UL79;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABI
(FRAGMENT).
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NON_TER
SEQUENCE
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
WYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-98277139; PubMed-9614934; MEDLINE-98277139; PubMed-9614934;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clin. Immunol. Immunopathol. EMBL; AF035033; AAD56269.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
NON_TER 1 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AF035031; AAI
HSSP; P80362; 1WTL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fetus."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Myosin-reactive autoantibodies in rheumatic carditis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Young D.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (FRAGMENT).
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                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                      TDFTLTISS
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108 AA;
                                                                                                                                                                                                                                                                                                                                Conservative
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11501 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                       108
11834 MW;
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Primates;
                                                                                                                                                                                                                                                                                                                                                  90.7%;
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                                                                                                                                                                                                                                                                                                                          Score 39; DB Pred. No. 1.1; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                       9F9C5A92EBA96EEA CRC64;
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                               CHAIN VARIABLE
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1.1;
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Best Local S
Matches
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                             InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF000047; Ig; 1.
SMART; SM00406; IGv; 1.
NON_TER 1 1 1
NON_TER 109 109
SEQUENCE 109 AA; 11928 MW;
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Q9UL86; Q9UL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (1)
SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Clin. Immunol. Immunopathol. EMBL; AF035028; AAD56264.1; HSSP; P80362; IWTL.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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"Myosin-reactive autoantibodies in rheumatic carditis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606; [1]
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Myosin-reactive autoantibodies in rheumatic carditis
70
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Q9UL7B;
Q9UL7B;
Q9UL7B;
Q9UL7B;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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MEDLINE-98277139; PubMed=9614934;
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Q9UL85;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN KAPPA CHAIN VARIABLE
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                          Clin. Immunol. Immunopathol. EMBL; AF035036; AAD56272.1;
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                                                                                                              SMART;
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                                                                                                                                                                                                                                                                                                                                           fetus."
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                                                                                                                                                                                                                                                                                                                                                                                                                 Young D.C.;
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                                                                                                    rPro; IPR003006; Ig_MHC.

rPro; IPR003596; Ig_v.

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T; SM00406; IGV; 1.
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                                                                                                                                                                                                                                              P80362;
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Pr00047; ig; 1.
; SM00406; IGV; 1.
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IPR003596; Ig_v.
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   109
11646 MW;
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Primates;
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Pred.
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          5F675C52EC7EE197 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                              rheumatic carditis
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Best Local
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EMBL; AE006628; AAK34738.1; -.

InterPro; IPR00559; FTHFS.

Pfam; PF01268; FTHFS: 1.F

PROSITE; PS00722; FTHFS_2; 1.

Ligase; Complete proteome.

SEQUENCE 557 AA; 59053 MW;
                                                                                                                                                                                                                  Q9JL74;
Q9JL74;
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=SF370 / ATCC 700294 / SEROTYPE M1;
MEDLINE=21192684; PubMed=11296296;
Ferretti J.J., McShan W.M., Ajdic D.J., Savic D.
Primeaux C., Sezate S., Suvorov A.N., Kenton S.;
Qian Y., Jia H.G., Najar F.Z., Ren Q., Zhu H., Y.;
Yuan X., Clifton S.W., Roe B.A., McLaughlin R.;
"Complete genome sequence of an M1 strain of Sty
Proc. Natl. Acad. Sci. U.S.A. 98:4658-4663(2001)
MEDLINE-204/8942; PubMed-10992488; Malkiel S., Liao L., Cunningham M.W., Diamond B.; "T-Cell-dependent antibody response to the dominant streptococcal polysaccharide, N-acetyl-glucosamine, with cardiac myosin."; Infect. Immun. 68:5803-5808(2000). EMBL; AF206032; AAF69330.1; -. HSSP; P80362; IWTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-DEC-2001 (TREMBLIEL 17, Last sequence update)
PUTATIVE FORMATE-TETRAHYDROFOLATE LIGASE (EC 6.3.4.3)
FHS.2 OR SPY2085.
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Q99xR2;
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Bacteria; Firmicutes;
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                                                                                         SEQUENCE FROM N.A. STRAIN-BALB/C;
                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. STRAIN-SF370 / ATC
                                                                                                                                                            Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Streptococcus
                                                                                                                           NCBI_TaxID=10090;
                                                                                                                                                                         (FRAGMENT)
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nicutes; Bacillus/Clostridium
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                                                                                                                                      Chordata;
Rodentia;
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LIGHT CHAIN VARIABLE REG
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                                                                                                                                     Craniata; Vertebrata; Sciurognathi; Muridae;
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Q96PF6;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
KAPPA 1 LIGHT CHAIN VARIABLE REGION (FRAGMENT).
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Q929J3;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
PUTATIVE PEPTIDOGLYCAN BOUND PROTEIN (LPXTG MOTIF
                                                                                                                                                                                                                                LIN2282.
Listeria innocua.
Bacteria; Firmicutes; Ba
Bacillus/Staphylococcus
      PubMed-11679669;
Glaser P., Frangeul L.,
Baquero F., Berche P.,
                                                                                                  SEQUENCE FROM N.A.
STRAIN-CLIP 11262 / SEROVAR 6A;
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MEDLINE-21361171; PubMed-11468171;
Comenzo R.L., Zhang Y., Martinez C., Osman K., Herrera G.A.;

"The tropism of organ involvement in primary systemic amyloidosis:
contributions of Ig V(L) germ line gene use and clonal plasma cell
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Homo sapiens (Human).
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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12735 MW;
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77.8%;
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   Bloecker H.,
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C., Amend A. Chakraborty
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                                                                                                                                                                                                                                                                                                                                     "Recombination in human herpesvirus 8 strains evolution of the K15 gene.";
Submitted (JUN-2001) to the EMBL/GenBank/DDBJ (EMBL; AY042942; AAK72676.1; -...
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Domann E., Dominguez-Bernal G., Duchaud E., Durant L., Dussurget O.,
Entian K.-D., Fsihi H., Garcia-del Portillo F., Garrido P.,
Gautier L., Goebel W., Gomez-Lopez N., Hain T., Hauf J., Jackson D.,
Jones L.-M., Kaerst U., Kreft J., Kuhn M., Kunst F., Kurapkat G.,
Madueno E., Maitournam A., Mata Vicente J., Ng E., Nedjari H.,
Nordsiek G., Novella S., de Pablos B., Perez-Diaz J.-C., Purcell R.,
Remmel B., Rose M., Schlueter T., Simoes N., Tierrez A.,
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"Comparative genomics of Listeria species.";
Science 294:849-852(2001).
EMBL; AL596171; CAC97510.1;
-. Listilist; LIN02282;
-. Lincoln Lindow Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kakoola D.N., Sheldon J., Byabazaire N., Bowden Katongole-Mbidde E., Schulz T.F., Davison A.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kaposi's sarcoma-associated herpesvirus (KSHV) (Human herpesvirus 8). Viruses; dsDNA viruses, no RNA stage; Herpesviridae; Gammaherpesvirinae; Rhadinovirus.
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on update)
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2: \SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1981.DAT: *
3: \SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1982.DAT: *
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6: \SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1985.DAT: *
7: \SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1987.DAT: *
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Gapop 10.0 , Gapext
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Copyright (c) 1993 - 2000 Compugen
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AAY52745
AAB98286
AAB98288
AAB97666
AAB97668
AAW80805
AAW809811
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AAW62807
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AAW52241	223	AAW52237	223	AAW52243	AAR88717	AAR98497	AAR98495	AAW01268	AAR54310	AAY98246	AAY95137	AAW62803	AAW01285	AAR54318	AAG93599	AAY56683	AAY34316	AAR72060	AAR72059	AAR72058	AAR72065	AAR72064	AAR72063	90	206	751	AAW62808	9	665	65	9	AAW59619	AAW62806
	<u>-151</u>	Ÿ	LD2-11-V	TD1/2-	an antibody ka	2E3 light cha	o 2F2 light cha	gion of HI	db AIH	qp120	Anti-qp120 antibod	Ω	gion of HI	HIV qp120 in			antibody 2.	.16 VK-1 L ch	.3 VK-1 L c	VK reg	434 VK-1 rec	K.17 VK-1 L	K.19 VK-1 L	.9 VK-1 L c	₽-	chain	no acid sequen	l peptide f	tial peptide f	tial pept	tial peptide f	i-RSV F p	Amino acid sequenc

ALIGNMENTS

AAR87043

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AAR87043 standard; Peptide;

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Anti-human IL-5 recombinant antibody - useful for preventing or reducing eosinophilia and for treating certain allergic diseases,
                       WPI; 1996-058412/06.
                                    Athwal DS,
                                               (CLLT ) CELLTECH THERAPEUTICS LTD
                                                            17-JUN-1994;
                                                                        16-JUN-1995;
                                                                                     28-DEC-1995.
                                                                                                 WO9535375-A1
                                                                                                            Homo sapiens
                                   Bodmer
                                                           94GB-0012230.
                                                                       95WO-GB01411.
                                   WW,
                                  Emtage
                                  JS;
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Humanised antibody; interleukin-5; IL-5; recombinant antibody; antibody engineering; monoclomal antibody; MAD; 39D10; CDR; complementarity determining region; light chain; framework; eosinophilia; allergy; asthma.

Human group I light chain framework 3.

25-JUN-1996 AAR87043;

(first entry)

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RESULT AAYSULT AAYSULT AAYSULT AAYSULT AAYSULT AAYS XX AAYS XX AAYS XX Huma KW AIR: KW
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The present invention describes chimeric antibody (Ab) heavy (H) chains CC containing the variable region of the H chain of a mouse monoclonal Ab CC recognising human tissue factor (hTF) and the constant region of the H CC chain of a human Ab. The variable region is one of six specified CC sequences (which are the H chain variable regions from mouse monoclonal CC Ab's ATR-2,3,4,5,7 or 8). Also described are chimeric Ab light (L) CC chains containing the variable region of the L chain of a mouse monoclonal Ab recognising human tissue factor (hTF) and the constant CC region of the L chain of a human Ab, the variable regions from mouse monoclonal Ab recognising human tissue factor (hTF) and the constant CC region of the L chain of a human Ab, the variable regions from mouse monoclonal Ab's ATR-2,3,4,5,7 or 8). The chain variable regions from mouse the treatment and prevention of thrombotic disease, especially of disseminated intravascular coagulation (DIC). The humanised antibody has the high'hTF binding activity of the mouse monoclonal antibody but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human tissue factor; TF; humanised; antibody; mouse monoclonal antibody; ATR-2; ATR-3; ATR-4; ATR-5; ATR-7; ATR-8; thrombotic disease; DIC;
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                                                                                                                                                                                                                                                                                                  (SLOK )
(SCRI )
                                           The present invention describes a method for treating cancers that express the A33 antigen. The method comprises administering an antigen antigen. The method comprises administering an immunoglobulin product (II) that binds specifically to A33 and contains one or more of 13 specified complementarity determining regions (CDRs), given in AAB98262 to AAB98274. (I) has cytostatic activity. The method can be used for treating colon and stomach cancers. (II), or the nucleic acid encoding it, can be used directly, in unconjugated form, for immunotherapy of cancer, and, when labeled, for detection or diagnosis of diseases associated with A33 expression. AAH22218 to AAH22244 and AAB98230 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB98286;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           greatly reduced immunogenicity. AAZ33001 to AAZ33091 and Y527007 to AAY52767 represent sequences used in the exemplification of the present
                                                                                                                                                                              Claim 16; Page 40; 85pp; English.
                                                                                                                                                                                                                    antigen by
                                                                                                                                                                                                                              Treating cancers,
                                                                                                                                                                                                                                                                           Barbas CF,
                                                                                                                                                                                                                                                                                                                                                  22-OCT-1999;
04-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; rabbit; humanised; A33 antigen; anti-A33 antigen antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-A33 antigen immunoglobulin
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Sequence
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                                    AAB98321 represent
                                                                                                                                                                                                       immunogiobulin product
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SCRIPPS RES INST.
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                                   sequences used
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Query Match Best Local Similarity

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RESULT
AAB98288
ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; rabbit; humanised; A33 antigen; anti-A33 antigen antibody; immunoglobulin; complementarity determining region; CDR; cancer; cytostatic; anticancer; colon cancer; stomach cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating cancers, particularly of stomach and colon, that express A33 antigen by administering conjugate of anticancer agent with specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200130393-A2
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                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                               present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Barbas CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (LUDW-) LUDWIG INST CANCER RES.
(SLOK ) SLOAN KETTERING INST CANCER RES
(SCRI ) SCRIPPS RES INST.
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04-APR-2000; 2000US-0543004.
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                                                                                                                                                                                                                                                                                                                                                                                       mammalian antibodies, which can be used for the treatment of a variety of diseases. The present sequence represents an A33 antigen binding immunoglobulin product VLFR3 peptide which is given in an example from
                                    Chimeric antibody; humanised; humanisation; mammalian antibody; immunoreact; anti A33 antigen antibody; immunoglobulin.
                                                                         A33 antigen binding immunoglobulin product VLFR3 peptide SEQ ID NO:94.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Preparing humanized rabbit antibodies that specifically immunoreact with a particular antigen using display technology for expressing libraries of antibody domains and fine tuning variable domain regions
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                                                                                                                                  AAB97668;
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                                                                                                                                                                                                                                                                               100.0%; ilarity 100.0%; Conservative
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Homo sapiens

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ARESULT AAW62805 ID AAW62805 ID AAW62805 ID AAW6 AC AAW6 XX AAW6 IDE AM11 XX IDE AM11 XX IDE AW110 IDE AW1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amino acid sequence of a human antibody fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           with a particular antigen using display technology for expressing libraries of antibody domains and fine tuning variable domain regions
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                       (ABGE-) ABGENIX INC
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                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                     Human; immunoglobulin; Ig; transgenic; non-human mammal;
inactivated endogenous Ig locus; B-ceil development;
human heavy chain Ig locus; micro constant region; J-Kappa gene;
kappa light chain Ig locus; kappa constant region; J-kappa gene;
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nes 9; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 77; 128pp; English.
                         WPI; 1999-034712/03.
                                                                                                           05-MAY-1997;
                                                                                                                                    05-MAY-1998;
                                                                                                                                                                                                                   Homo sapiens
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                                                                              (ABGE-) ABGENIX
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                                                                                                                                                                                                                                                            epidermal growth factor receptor; tumour; EGF
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Humanised antibodies against epidermal growth factor receptor, EGF-r

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RESULT AAW62807
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AAW62793-822 represent fragments of human antibodies produced by transgenic xenomice, created using the method of the invention. 'specification describes a transgenic non-human mammal which has a modifications that comprise an inactivated endogenous immunoglob
                                                                                                                                                                                                                                                       New transgenic non-human mammals - having immunoglobulin locus and a near complete ! used for production of human antibodies
                                                                                                                                                                                          Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; immunoglobulin; Ig;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ABGE-) ABGENIX INC
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which has genome immunoglobulin (
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Matches 9
            AAW14490-91 are the heavy and light chains (respectively) of monoclonal antibody (MAb) D. MAb D recognises Vibrio cholera neuraminidase-resistant epitope of ganglioside GD2, from a human melanoma cell line. MAbs A, B and C (see AAW14448-98) are mentioned in the specification, but are not part of the claims. MAbs A and B recognise antigens 3 and 11 resp., of a permenant human tumour cell line. MAb C also recognises an epitope of a tumour-associated antigen. These antigens
                                                                                                                                                                                                                                                              Disclosure; Page 14; 19pp; German.
                                                                                                                                                                                                                                                                                                                    gastrointestinal tumour marker
                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; AAT63508
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                                                                                                                                                                                                                                                                                                                                                   Monoclonal antibody to tumour-associated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-MAR-1989;
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   concns.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monoclonal antibody D VK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR99878 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumours, e.g. pancreatic carcinoma, and are thus useful as tumour markers for diagnostic or therapeutic purposes.
                                                                                                                                                          tumour cell line.
MAD D (AAT35665-T36666) recognises a Vibrio cholera neuraminidase-
resistant epitope of ganglioside GD2, from a human melanoma cell
                                                                                                                                                                                                                                            MADS A, B and D are mentioned in the specification, but are not part of the claims.

MAD A (AAT36661-T36662) recognises antigen 3 of permenant human tumour cell line.
                                                                                                                                                                                                                                                                                                                                 MAb C (AAT36659-T36660) is a monoclonal antibody that recognises an epitope of a tumour-associated antigen occurring at high concn. in the serum of patients with gastrointestinal tumours, e.g. pancreatic carcinoma, and is thus useful as a tumour marker for diagnostic or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-372835/38.
N-PSDB; AAT36666.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 14; 19pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          gastrointestinal tumour marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Auerbach B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-AUG-1996
                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibody to tumour-associated antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BEHW ) BEHRINGWERKE AG
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                                                                                                                                                                                                                              MAD B (AAT36663-T36664)
                                                                                                                                                                                                                                                                                                                  therapeutic purposes.
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hes 9; Conserv
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llarity 100.0%;
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                                  Conservative
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                                                                                                                                                                                                                            recognises antigen 11 of permenant human
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Pred. No.
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0.36;
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cc transgenic Xenomice, created using the method of the invention. The modification describes a transgenic non-human mammal which has genome cc modifications that comprise an inactivated endogenous immunoglobulin (Ig) cc locus, so that the mammal does not display normal B-cell development. The modified genome also has an inserted human heavy chain Ig locus in cc germline configuration, the human heavy chain Ig locus comprising a human cc micro constant region and regulatory and switch sequences, human J-H cc genes, human D-H genes, and human V-H genes and an inserted human kappa light chain Ig locus comprising a human V-H genes and an inserted human kappa genes, where the number of V-H and V-kappa genes inserted cc and V-kappa genes, where the number of V-H and V-kappa genes inserted cc are selected to restore normal B-cell development in the mammal. The cc transgenic animals have a near complete human Ig locus, including both a cc bused for the production of human antibodies when exposed to particular antigens e.g. when exposed to human II-B, EGFR or TNF- alpha cr restorively
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                                                                       Query Match
Best Local
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Mendez M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H kappa light chain Ig locus; kappa constant region; J-kappa gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW62793-822 represent fragments of human antibodies produced by transgenic Xenomice, created using the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   production; antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amino acid sequence of a human antibody fragment
                                                                                                                                                                                   Sequence
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                                             100.0%;
nilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  86
                                                   0;
                                                   Score 43; DE Pred. No. 0.3); Mismatches
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                                                                               DB 19;
. 0.37;
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V-kappa;
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RESULT 1
AAY56654
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                                                                                                          Query Match
Best Local Similarity
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AAW59619
                                                                                                                                                                                                                 Dpk9 antibody. The invention relates to the construction and use of human MAbs specific for the fusion (F) protein of respiratory syncytial virus (RSV) to passively treat, prevent or detect RSV infection. Hul9A, Hul9B, Hul9C and Hul9D MAbs are claimed. These are reshaped human antibodies comprising a heavy chain selected from 19A, 19B, 19C or 19D (see AAM59615-18), which are based on the GL Dp58 sequence, and a light chain selected from 19A, 19B, 19C or 19D (see AAM59615-18), which are neutralising; they inhibit virus growth in vitro and in vivo in animal models of RSV infection. Nucleic acids encoding the human MAbs, recombinant plasmids (see AAW41427-33) and host cells (e.g. COS, CHO, myeloma)
          AAY56654 standard;
                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                          This is the heavy chain amino acid sequence of germline
                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibodies reactive with Respiratory Syncytial Virus useful for detection, prevention and treatment of RSV infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-286600/25.
                                                                                                                                                                                                          are provided.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monoclonal antibody; human; Hu19; engineered antibody; RSV; respiratory syncytial virus; fusion protein; infection; complementarity determining region; CDR; therapy; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9819704-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW59619;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW59619 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Region
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                                                                                                                    100.0%;
ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label- CDR1
50..57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /label= CDR2
         protein; 88
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                                                                                                                                Score 43; DB 1
Pred. No. 0.38;
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                                                                                                                       Mismatches
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                                                                                                                      Indels
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RESULT
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                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Thes 9; Conservative
                                                                       Complementarity determining region; antibody; primate; immunogenicity; old World ape; Old World monkey; antigen-binding affinity.
                                                                                                                                                                                                                                                                                                                                                          complementarity determining regions) derived from a non-human antigen-specific donor antibody, and an acceptor framework from a non-human primate. The Abs are prepared by grafting CDRs from a non-human antigen-specific donor antibody onto homologous old World ape or monkey acceptor frameworks. The Abs have reduced immunogenicity and are better tolerated in humans (because of the close similarity between the human and primate proteins), but retain the full antigen-binding affinity of the donor antibody.
                               WO9955369-A1
                                                 Pan troglodytes
                                                                                                       Partial peptide fragment of chimpanzee Vkappa cDNA clone 46-6
                                                                                                                                  15-FEB-2000
                                                                                                                                                        AAY56655;
                                                                                                                                                                            AAY56655 standard;
                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 22; Page 70-71; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antibodies containing donor complementarity determining regions and non-human primate acceptor frameworks, having reduced immunogenicity humans -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-023265/02.
N-PSDB; AAZ39322.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Taylor AH;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Complementarity determining region; antibody; primate; immunogenicity; Old World ape; Old World monkey; antigen-binding affinity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-APR-1998;
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                                                                                                                                                                                                                                                                               Score 43; DE
Pred. No. 0.3
0; Mismatches
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                                                                                                                                                                                                                                                                                         DB 2
0.38;
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Search completed: July 15, 2002, 12:57:58 Job time: 414 sec
                                                                        В
                                                                                                                            Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 9; Conservative 0;
                                                                                                                                                                                                                              The invention provides an antibody (Ab) comprising donor CDRs (complementarity determining regions) derived from a non-human antigen-specific donor antibody, and an acceptor framework from a non-human primate. The Abs are prepared by grafting CDRs from a non-human antigen-specific donor antibody onto homologous Old World ape or monkey acceptor frameworks. The Abs have reduced immunogenicity and are better tolerated in humans (because of the close similarity between the human and primate proteins), but retain the full antigen-binding affinity of the donor antibody.
                                                                                                                                                                                                                                                                                                                                                                                                             Antibodies containing donor complementarity determining regions and non-human primate acceptor frameworks, having reduced immunogenicity in humans -
                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                              Claim 22; Page 71; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-023265/02.
N-PSDB; AAZ39323.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Taylor AH;
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Maximum Match 100%
Listing first 45 summaries
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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seq length: 2000000000
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1: /cgn2_6/ptodata/2,
2: /cgn2_6/ptodata/2,
3: /cgn2_6/ptodata/2,
4: /cgn2_6/ptodata/2,
5: /cgn2_6/ptodata/2,
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2.905 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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PCT-US91-02942-42
US-08-470-139-13
US-08-569-147-40
US-09-247-061-13
US-09-245-638A-92
US-09-243-004-94
US-09-543-004-92
US-09-543-004-92
US-08-765-179B-10
US-08-765-179B-10
US-08-765-179B-14
US-08-554-840-12
US-08-851-362D-21
US-08-851-362D-21
US-08-854-840-13
US-08-89-554-840-13
US-08-89-575-106
US-08-899-575-106
US-08-899-575-89
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Compugen Ltd
Sequence 42, Appl Sequence 13, Appl Sequence 94, Appl Sequence 94, Appl Sequence 94, Appl Sequence 97, Appl Sequence 97, Appl Sequence 97, Appl Sequence 10, Appl Sequence 11, Appl Sequence 12, Appl Sequence 12, Appl Sequence 13, Appl Sequence 15, Appl Sequence 15, Appl Sequence 15, Appl Sequence 16, Appl Sequence 16, Appl Sequence 106, Appl Sequence 106
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PCT-US91-02942-42
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                                                                           Query Match
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Matches 9
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CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                      LENGTH:
         1 TDFTLTISS 9
                                                                              Similarity
9; Conserv
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USA
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45	44	4 3	42	41	40	39	38	37	36	35	34	<u>3</u> 3	32	31	30	29	28
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107	107	107	107	107	107	107	106	106	106	106	106	106	106	106	106	106	106
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US-08-300-386A-66	US-08-276-852-105	US-08-276-852-104	US-08-276-852-84	US-08-276-852-82	US-08-477-728-103	US-07-634-278-103	PCT-US95-08743-85	PCT-US95-08743-83	US-09-240-274-165	US-09-240-274-159	US-08-397-411-8	US-08-899-575-85	US-08-899-575-83	US-08-899-575-85	US-08-899-575-83	US-08-276-852-85	US-08-276-852-83
Sequence 66, Appl	Sequence 105, App		Sequence 84, Appi	-		Sequence 103, App	Sequence 85, Appl				Sequence 8, Appli	Sequence 85, Appl	Sequence 83, Appl	•	Sequence 83, Appl	•	Sequence 83, Appl

ALIGNMENTS

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TOPOLOGY: linear;
MOLECULE TYPE: protein
PCT-US91-02942-42
                                                                                                                                                                                                                           TELEFAX: (202) 833-8716
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US91/02942
FILING DATE: 19910429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ROTHLEIN, ROBERT
APPLICANT: ADAIR, JOHN R
APPLICANT: ATHMAL DILIZET S
TITLE OF INVENTION: HUMANIZED CDR-GRAFTED ICAM-1 ANTIBODY
NUMBER OF SEQUENCES: 102
CORRESPONDENCE ADDRESS: Sterne, Kessler, Goldstein & Fox.
                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9009549.8
FILING DATE: 27-APR-1990
ATTORNEY/AGENT INFORMATION:
NAME: FOX, SAM L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 30,353
REFERENCE/DOCKET NUMBER: 1011.0586600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 466-0800
                                                                                                                                                                                     AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Ave. NW Suite 300
       Conservative
                          100.0%;
       0;
Score 43; DB 5; Length 25;
Pred. No. 0.058;
Nismatches 0; Indels
         0;
         Gaps
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US-08-470-139-13; Sequence 13, Ap; Patent No. 5998
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Best Local Similarity
Thomas 9; Conserv
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; TOPOLOGY: 11n
; MOLECULE TYPE:
US-08-470-139-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Vers

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/569,147

FILING DATE: 25-March-1996

CLASSIFICATION: 536

ATTORNEY/AGENT INFORMATION:

NAME: Trujillo, Doreen Yakko

REGISTRATION NUMBER: 35,719

REFERNCE/DOCKET NUMBER: CARP-0047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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TITLE OF IN.

NUMBER OF SEQUENCES.

NUMBER OF SEQUENCES.

COMPUTER READABLE FORM:

COMPUTER FLADABLE FORM:

FLADABLE FORM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: HUMANISED ANTIBODIES NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
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NAME: TRUJILLO, DOREEN YATKO
REGISTRATION NUMBER: 35,719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U
ZIP: 19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: One Liberty CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & ADDRESSEE: No. 6180377ris, LLP STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 TDFTLTISS 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acids
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6180377
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5998586
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JER: CARP-0044
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0.075;
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APPLICANT: Bodmer, Mark
APPLICANT: Athwal, Diljeet Singh
APPLICANT: Emtage, John Spencer
TITLE OF INVENTION: Interleukin-5
FILE REFERENCE: CAPP-^^~
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                                                               SEQ ID NO 92
LENGTH: 32
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3
SEO ID NO 13
LENGTH: 32
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 92, Application US/09425638A Patent No. 6342587
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                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/425,638A
CURRENT FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 129
                                                                                                                                                                                                                                                     APPLICANT: Carlos F. Barbas III, Christoph Rader, Gerd Ritter, Sydney Welt an APPLICANT: Lloyd J. Old
TITLE OF INVENTION: A33 ANTIGEN SPECIFIC IMMUNOGLOBULIN PRODUCTS AND USES THE FILE REFERENCE: LUD 5630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 9; Conserv
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CURRENT FILING DATE: 1999-07-02
NUMBER OF SEQ ID NOS: 28
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FEATURE:
                     ORGANISM: Homo sapiens
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nes 9; Conserv
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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JS-09-425-638A-92

Patent No. 6346249

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Query Match
Best Local Similarity
Whiches 9; Conserv
RESULT 8
US-09-543-004-94
; Sequence 94, Application US/09543004
                                                                                                                       Query Match
Best Local Similarity
"""+"hes 9; Conserv
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Best Local Similarity
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US-09-425-638A-94
                                                                                                                                                                                                                                                     SEQ ID NO 92
LENGTH: 32
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 94
LENGTH: 32
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 92, Application US/09543004 Patent No. 6346249
                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/543,004
CURRENT FILING DATE: 2000-04-04
PRIOR APPLICATION NUMBER: 09/425,638
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/425,638A CURRENT FILING DATE: 1999-10-22
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TITLE OF INVENTION: A33 ANTIGEN SPECIFIC IMMUNOGLOBULIN PRODUCTS AND USES THER
                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Carlos F. Barbas III, Christoph Rader, Gerd Ritter, Sydney Welt and APPLICANT: Lloyd J. Old
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                            FEATURE:
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                                                                              IDFILIISS 21
                                                                                                                                    100.0%;
ilarity 100.0%;
Conservative 0;
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Pred. No. 0.075;
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                                                                                                                                                   Score 43; DB 4;
Pred. No. 0.075;
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; LENGTH: 32
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; FEATURE:
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Best Local Similarity
"hes 9; Conserva
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; TOPOLOGY: linear; MOLECULE TYPE: peptide PCT-US91-02942-9
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APPLICANT: Carlos F.
APPLICANT: Lloyd J. Old
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/543,004
CURRENT FILING DATE: 2000-04-04
PRIOR APPLICATION NUMBER: 09/425,638
PRIOR FILING DATE: 1999-10-22
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                                                                                                  TELEFAX: (202) 833-87
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9009549.8
FILING DATE: 27-APR-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy
                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ATHWAL, DILJEET S
TITLE OF INVENTION: HUMANIZED CDR-GRAFTED ICAM-1 ANTIBODY
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PEILING DATE: 19910429 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Washington STATE: D.C.
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                                              AMINO ACID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ROTHLEIN, ROBERT
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                                                                                                                 (202)
                                                                                                                                 (202) 466-0800
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APPLICATION NUMBER: PCT/EP95/02626
FILING DATE: 06-JUL-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 44 25 115.7
FILING DATE: 15-JUL-1994
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
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Best Local Similarity
"atches 9; Conservi
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                                                                                                                                                                     Sequence 14,
Patent No. 5
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                                                                                                                                                       GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,179B
FILING DATE: 14-JAN-1997
                                                                                 APPLICANT: STEIPE, Boris
APPLICANT: STEINBACHER, Stefan
TITLE OF INVENTION: PROCESS FOR MODIFYING
TITLE OF INVENTION: OF ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 28

CORRESPONDENCE ADDRESS:
ADDRESSE: NIKaido, Marmelstein, Murray & Oram LLP
STREET: 655 Flfteenth Street N.W. Suite 330
                                                 NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 11
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                   STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
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                               ADDRESSEE:
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20005-5701
Washington
                                                                                                                                                                         1, Application US/08765179B 5854027
               E: Nikaido, Marmelstein, Murray & Oram LLP 655 Fifteenth Street N.W. Suite 330
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Pred. No. 0.16;
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; MOLECULE TYPE: protein
US-08-765-179B-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9, Application US/08554840 Patent No. 6001358
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APPLICANT: BLACK, Amelia
APPLICANT: HANNA, Nabil
APPLICANT: PADLAN, Eduardo A.
APPLICANT: NEWMAN, Roland A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: DE P 44 25 115.7 ETIJING DATE: 15-UUL-1994 INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS: LENGTH: 64 amino acids
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,
                            TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 2: CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/02626
FILING DATE: 06-JUL-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                        STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
                                                                                                                                                  APPLICATION NUMBER: US/08/554,840 FILING DATE: 07-NOV-1995
                                                                                                                                                                                                                                                                                      STATE: Virginia COUNTRY: United States ZIP: 22313-1404
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nes 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                    CLASSIFICATION: 536
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                                                                                                                                                                                                                                                                                                                                                       E: Burns, Doane, Swecker & Mathis P.O. Box 1404
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                 (703)
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HUMANIZED ANTIBODIES TO HUMAN 9p39,

COMPOSITIONS CONTAINING AND THERAPEUTIC USE THEREOF
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Pred. No. 0.16;
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Length 64; Indels

SEQUENCE CHARACTERISTICS:

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APPLICANT: Jakobovits, Aya
APPLICANT: Jakobovits, Aya
APPLICANT: Yang, Xiao-Dong
APPLICANT: Gallo, Michael
APPLICANT: Jia, Xiao-Chi
FITLE OF INVENTION: Growth Factor Receptor
FILE OF INVENTION: Growth Factor Receptor
FILE REFERENCE: Cell 4.20
CURRENT APPLICATION NUMBER: US/08/851,362D
CURRENT FILIKG DATE: 1997-05-05
NUMBER OF SEQ ID NOS: 49
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 76
TYPE: PRT
ORGANISM: human
US-08-851-362D-21
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Best Local Similarity 100.0%;
Matches 9; Conservative 0;
                                                                                                                                                                                                                                                                                               Sequence 12, Application US/08554840 Patent No. 6001358
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                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                APPLICANT: PADLAN,
APPLICANT: NEWMAN,
                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                 APPLICANT:
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TOPOLOGY: lin
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                                                          COUNTRY: United ZIP: 22313-1404
                                                                                         STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
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                                                                           United States
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, Roland A.
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100.0%; Pred. No. 0.19;
PC-DOS/MS-DOS
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Pred. No. 0.17;
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; MOLECULE TYPE:
US-08-554-840-12
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INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 80 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 9
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Patent No. 6
                                                                                                                                   COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/554,840
FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 012712-127
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
TELEPHONE: (703) 836-6620
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INFORMATION FOR SEQ ID NO:
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APPLICANT:
APPLICANT:
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NAME: Teskin, Robin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: HUNTITLE OF INVENTION: CON NUMBER OF SEQUENCES: 28 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-6620
                                                                                    SEQUENCE CHARACTERISTICS:
TOPOLOGY: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Teskin, Robin L. REGISTRATION NUMBER: 35,030
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/0
FILING DATE: 07-NOV-1995
                               LENGTH: 80 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: United States ZIP: 22313-1404
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                                                                                                                        (703) 836-2021
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Nabil
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Maximum Match 10
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 protein search, using sw model
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Match
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58
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ALIGNMENTS

trans-activating transcription regulator - human immunodeficiency virus type 1 (isola C.Species: human immunodeficiency virus type 1, HIV-1 A:Note: host Homo sapiens (man)
C.Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 16-Jul-1999
C.Accession: JQ0071
R.Spire, B.; Sire, J.; Zachar, V.; Rey, F.; Barre-Sinoussi, F.; Galibert, F.; Hampe, Gene 81, 275-284, 1989
A:Title: Nucleotide sequence of HIVI-NDK: a highly cytopathic strain of the human immunity. A:Reference number: JQ0065; MUID:90034200 A;Molecule type: DNA A;Residues: 1-86 <SPI> A;Cross-references: GB:M27323; NID:g328154; PIDN:AAA44866.1; PID:g328155 RESULT TNLJND C; Genetics: A;Gene: tat

C;Superfamily: AIDS trans-activating transcription
C;Keywords: AIDS; immunodeficiency; transcription regulator

Ş Query Match Best Local S Matches 9 Local Similarity 81.1 hes 9; Conservative 1 YGRKKRRQKKK 11 89.7%; 81.8%; 2; Score 52; DB 1 Pred. No. 0.15; Mismatches DB 1; 0; Length 86; Indels 0;

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В

47

YGRKKRRORRK 57

A; Gene: tat C; Superfamily: 1 C; Keywords: trai trans-activating transcription regulator - human immunodeficiency virus type 1 c;Species: human immunodeficiency virus type 1, HIV-1 C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 21-Jul-2000 C;Accession: T09384 A;Molecule type: DNA
A;Residues: 1-71 <MIC>
A;Cross-references: EMBL:U24451; NID:g829440; PIDN:AAA79576.1; PID:g829444
C;Genetics: R.Michael, N.L.; Chang, G.; d'ARCY, L.A.; Ehrenberg, P.K.; Mariani, R.; Busch, M.P.; J. Virol. 69, 4228-4236, 1995
A.Title: Defective accessory genes in a human immunodeficiency virus type 1-infected A;Reference number: Z16654; MUID:95287475
A;Accession: T99384
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA ν (isola

Query Match

84.5%; Score 49; DВ 2; Length 71; ly: AIDS trans-activating transcription regulator transcription

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RESULT
A25700
trans-activating transcription regulator - human immunodeficiency virus type C;Species: human immunodeficiency virus type 1, HIV-1 C;Date: 28-Sep-1987 #sequence_revision 28-Sep-1987 #text_change 02-Jul-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        trans-activating transcription regulator - human immunodeficiency virus C:Species: human immunodeficiency virus Er-6 C:Date: 30-Sep-1987 #sequence_revision 30-Sep-1987 #text_change 16-Jul-1 C:Accession: C26192 R:Srinivasan, A.; Anand, R.; York, D.; Ranganathan, P.; Feorino, P.; Scherick, A.; Anand, R.; York, D.; Ranganathan, P.; Feorino, P.; Scherick, A.; Anand, R.; York, D.; Ranganathan, P.; Feorino, P.; Scherick, A.; Anand, R.; York, D.; Ranganathan, P.; Feorino, P.; Scherick, A.; Anand, R.; York, D.; Ranganathan, P.; Feorino, P.; Scherick, A.; Anand, R.; York, D.; Ranganathan, P.; Feorino, P.; Scherick, R.; Feorino, P.; Feorino, P.; Scherick, R.; Feorino, P.; Scherick, R.; Feorino, P.; Feorino, P.; Feorino, P.; Scherick, R.; Feorino, P.; 
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R;Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, V.S.; Casey, J.M.; Srinivasan, A · Anderson D · Desai, S.M.; Kalyanaraman, A · Anderson D · Desai, S.M.; Kalyanaraman, A · Desai, S.
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C; Superfamily: /
C; Keywords: AID:
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A:Title: Molecular characterization of human
A:Reference number: A26192; MUID:87248097
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A:Gene: tat
A:Gene: tat
C:Superfamily: AIDS trans-activating
C:Keywords: transcription regulation
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A; Residues: 1-86 <SRI>
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A; Residues: 1-72 < DES>
A; Cross-references: GB: M13137; NID: g326460
A; Note: the GenBank entry ADRE3AA PID: g209908
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A:Title: Molecular cloning and primary nucleotide sequence & A;Reference number: A94136; MUID:87041461
A;Accession: B25523
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3; Mismatc
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A; Molecule type: nucleic acid
A; Residues: 1-86 <SID>
A; Cross references: EMBL:X64650
C; Genetics:
A; Gene: tat
A; Introns: 72/2
C; Superfamily: AIDS trans-activ
                                                                                                                                                                                                                                                                                                                                 trans-activating transcription regulator - human immu c;Species: human immunodeficiency virus type 1, HIV-1 c;Date: 06-Oct-1994 #sequence_revision 23-Feb-1996 #tC;Accession: S33982; S26385; S19864 R;Carilni, F.
                                                                                                                           A;Cross-references: EMBL:Z11530; NID:g60192; PIDN:CAA77625.1; PID:g60196
R;Siderovski, D.P.; Matsuyama, T.; Frigerio, E.; Chui, S.; Min, X.; Erfle, H.; Sumner Nucleic Acids Res. 20, 5311-5320, 1992
Nucleic Acids Res. 20, 5311-5320, 1992
A;Title: Random mutagenesis of the human immunodeficiency virus type-1 trans-activato A;Reference number: S26385; MUID:93065196
A;Accession: S26385
                                                                                                                                                                                                                                        A; Reference number: S3
A; Accession: S33982
A; Molecule type: DNA
A; Residues: 1-86 <CAR>
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S33982
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C;Species: human immunodeficiency virus type 1, HIV
C;Date: 15-Jul-1995 #sequence_revision 01-Sep-1995
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S54381
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A; Status: preliminary
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A;Residues: 1-86 <SOD>
C;Superfamily: AIDS trans-activating transcription
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R;Sodroski, J.; Patarca, R.;
Science 229, 74-77, 1985
Science 229, 74-77, 1985
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                                                                          NID: g60144;
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Query Match
Best Local Similarity
----hes 8; Conserv
                                                                                                                                                                                                                                                                                                                                         tat protein - human immunodeficiency virus type 1
(;Species: human immunodeficiency virus type 1, HIV-1
C;Date: 19-Feb-1999 #sequence_revision 19-Feb-1999 #text_change
C;Accession: T01665
R;Alizon, M.: Wain-Hobson, S.: Gluckman, J.C.; Sonigo, P.
Cell 46, 63-74, 1986
A;Title: Genetic variability of the AIDS virus: Nucleotide seque
A;Reference number: Z14389; MUID:86245056
    trans-activating
C; Species: human
                                                                                                                        Qy
                                                                                                                                                                                                                      A;Introns: 72/2
C;Superfamily: AIDS trans-activating transcription regulator
                                     TNLJ12
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A; Residues: 1-87 <ALI>
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Best Local Similarity
"hes 8; Conserv:
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A:Title: A selection system to study protein-RNA interactions: Functional display A:Reference number: JC5591; MUID:97350867
A:Accession: JC5591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N;Alternate names: tat protein
C;Specles: human immunodeficiency virus type 1, HIV
C;Date: 23-Sep-1997 #sequence_revision 23-Sep-1997
C;Accession: JC5591
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R; Hoffmann, S.; Willbold, D.
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8; Conservative
transcription regulator - human immunodeficiency virus type 1 (isolate immunodeficiency virus type 1, HIV-1 \,
                                                                                                                                                          Conservative
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tat protein - human immunodeficiency virus type 1 (strain JRFL) c;Species: human immunodeficiency virus type 1, HIV-1 C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change C;Accession: T09446
R;Pang, S.; Vinters, H.V.; Akashi, T.; O'Brien, W.A.; Chen, I.S.
A;Cross-references: EMBL:U63632; NID:g1465777; PID:g1465783 C;Genetics: A;Gene: tat
                                                                                                                                                                                                                    submitted to the EMBL Data A; Reference number: 216673 A; Accession: T09446
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T09446
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                                                                                                        A; Molecule type: DNA
A; Residues: 1-101 < PAN>
                                                                                                                                                                             A; Status: preliminary; translated
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C;Superfamily: AII
C;Keywords: AIDS;
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A;Gene: .tat
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A; Residues: 1-101 <LIY>
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C;Superfamily: AIDS trans-activating
C;Keywords: AIDS; immunodeficiency;
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A;Title: Three novel genes of human T-lymphotropic virus type
A;Reference number: A94093; MUID:86177573
A;Accession: A04017
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A; Residues: 1-95
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72.7%;
                                                                                                                                                                                                                                                                                     Akashi, T.; O'Brien, W.A.;
Library, July 1996
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72.7%;
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3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   n regulator
regulation
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C; Accession: D38095; Sbl8//; Sbl800
C; Accession: D38095; MJD: B501-8536, 1992
J. Biol. Chem. 267, 8530-8536, 1992
A; Title: The human T cell transcription factor-1 gene. Structure, localization, and p A; Reterence number: A38095; MUID:9225082
A; Status: preliminary
A; Accession: D38095
A; Status: preliminary
A; Accession: Colored type: DNA
A; Cross references: G81877
A; Clevers, H.; Ballhausen, W.G.
Submitted to the EMBL Data Library, January 1995
A; Description: The human high mobility group (HMG)-box transcription factor TCF-1: no A; Reference number: S61877
A; Accession: S61877
A; Clevers, H.; Ballhausen, W.G.
Biochim. Biophys. Acta 1263, 169-172, 1995
A; Title: The human high mobility group (HMG)-box transcription factor TCF-1: novel is A; Reference number: S61796; MUID:95367594
A; Re
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C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change
C;Accession: D38095; S61877; S61800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ş
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C; Superfamily:
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R:van de' Wetering, M.; Oosterwegel, M.; Holstege, F.; I
J. Biol. Chem. 267, 8530-8536, 1992
A;Title: The human T cell transcription factor-1 gene.
A;Reference number: A38095; MUID:92235082
                                                                                                               N;Alternate names: transcription factor TCF-IC C;Speciles: Homo sapiens (man) C;Speciles: Homo sapiens (man) C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 24-Sep-1999 C;Date: 19-May-1994 #text_change 24-Sep-1999 C;Accession: C38095; S13449; S65007
                                                                                                                                                                                                      T-cell-specific transcription factor 1 splice form C - N; Alternate names: transcription factor TCF-1C
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
7; Conserva
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72.7%;
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63.6%;
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Pred. No. 0.52
3; Mismatches
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Pred. No. 9.7;
3; Mismatches
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                                                                                          F.; Dooyes, D.; Suijkerbuijk,
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                                    Structure,
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erences A38095 and S61796
R;Mayer, K; Wolff, E; Clevers, H; Ballhausen, Biochim. Biophys. Acta 1263, 169-172, 1995
A;Title: The human high mobility group (HMG)-box A;Reference number: S61796; MUID:95367594
A;Accession: S65007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBO J. 10, 123-132, 1991

A;Title: Identification and cloning of TCF-1, a T lymphocyte-specific transcription A;Reference number: $13449; MUID:91114695

A;Accession: $13449
                                                                                                                                                                                                                                                                                                                                                                                     A; Reference number: A38095; MUID:92235082

A; Accession: A38095

A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-269 <VAN>
A; Residues: 1269 <VAN>
A; Residues: GB:X63901; NID:g36791; PIDN:CAB56795.1; PID:g6006565
A; Note: authors translated the codon GAT for residue 253 as Ser
R; van de Wetering, M; Costerwegel, M; Dooijes, D; Clevers, H.
EMBO J. 10, 123-132, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T-cell-specific transcription factor 1 splice form A - human N,Alternate names: transcription factor TCF-1A C;Species: Homo sapiens (man) C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 21-Jul-2000 C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 21-Jul-2000 C;Accession: A38095; 816645; 865005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 244-268 <MAY>
A; Note: only a part of the
C; Genetics:
                                                                        A; Molecule type: mRNA
A; Residues: 1-269 
A; Residues: 1-269 
A; Clevers; H.; Ballhausen, W.G.
A; Clevers, K.; Wolff, E.; Clevers, H.; Ballhausen, W.G.
Blochim. Blophys. Acta 1263, 169-172, 1995
A; Title: The human high mobility group (HMG)-box transcription
A; Reference number: S61796; MUID: 95367594
A; A; A; Ccession: S65005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rivan de Wetering, M.; Oosterwegel, M.; Holstege, F.; J. Biol. Chem. 267, 8530-8536, 1992
A;Title: The human T cell transcription factor-1 gene. A;Reference number: A38095; MUID:92235082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Superfamily: unassigned HMG box proteins; HMG box homology C;Keywords: alternative splicing; DNA binding; transcription F;151-226/Domain: HMG box homology <HMGl>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R; van de Wetering, M.;
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A; Residues: 1-268 < VAN>
A; Status: not compared with A; Molecule type: mRNA A; Residues: 244-269 <MAY>
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                                                                                                                                                                                                                                                                                                         A; Reference number: A; Accession: S16645
                                                                                                                                                                                                                                                                                                                                      A; Title: Identification and cloning of TCF-1, a A; Reference number: S13449; MUID:91114695
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 7; Conserv
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                                                          conceptual
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Pred. No. 10;
3; Mismatches
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                                                             translation
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C:Generics:
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C:Supericmnly; unassigned HMG box proteins; HMG box homology
C:Supericmnly; unassigned HMG box proteins; HMG box homology
C:Box transcription factor
F:151-228/homain: HMG box homology <:HMClr

Query March
74.1%; Score 43; DB 2; Length 269;
HMCChes 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;
V 1 YGRKENDKEK 11
DB 227 YGKKENSEE 237

Search completed: July 15, 2002, 13:01:03

Job time: 469 sec
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P18804;
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-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIINE*90034200; PubMed*2806917;
Spire B., Sire J., Zachar V., Rey F., Barre-Sinoussi F., Galibert F.,
Hampe A., Chermann J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/anor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus type 1 (NDK isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus. NCBI_TaxID=11695;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1990
16-OCT-2001
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47
                                      1 YGRKKRRQKKK 11
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                                                                                                                                                                                      DB 1;
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                                                                                                                         0;
                                                                                                                                                                                      Length 86;
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RESULT
TAT_HV1B5
ID TAT_HV1B5
P04612;
P40G-10
      RESULT 3
TAT_HV112
ID TAT_H
AC P0432
AC 20-MA
DT 13-AU
DT 13-AU
DT TAT P
GN TAT P
GN TAT
OS Human
OX NCBI
RN [1]*.
RN [1]*.
RN SEQUE
RX MEDLI
RX ME
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Best Local S
Matches 8
                                                                                                                                                                                                          20-MAR-1987
13-AUG-1987
16-OCT-2001
                                                                                                                                                                                                                                                                           TAT_HV112
P04326;
    Arya
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13-AUG-1987
16-OCT-2001
                           MEDLINE-86177573;
                                              SEQUENCE FROM N.A.
                                                                                        Human immunodeficiency virus type 1 (clone 12) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus.
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for come entities requires a license agreement (See http://www.isb-sib.ch/anor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Complete nucleotide sequence of the AIDS virus, HTLV-III.";
Nature 313:277-284(1995).

-1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TTRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.

MEDILINE-85111123; PubMed-2578615;

Ratner L., Haseltine W., Patarca R., Livak K.J., Starcich B.R

Josephs S.F., Doran E.R., Rafalski J.A., Whitehorn E.A.,

Baumeister K., Ivanoff L., Petteway S.R. Jr., Pearson M.L.,

Lautenberger J.A., Papas T.S., Ghrayeb J., Chang N.T., Gallo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Viruses; Retroid
NCBI_TaxID-11682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus type 1 (BH5 isolate) (HIV-1).
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                                                                                                                                                                                                                                                                                                                                                                                                              47
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                                                                                                                                                                                protein
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S.K., Gallo R.
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8; Conser
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0539; Tat; 1.
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                                                                                                                                                                                                     (Rel. 04, Created)
(Rel. 05, Last seq
(Rel. 40, Last ann
                                                                                                                                                                            (Rel. 40, Last annotation update) (Transactivating regulatory protein).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
(Transactivating regulatory protein)
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8 AA;
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                         PubMed=3008154;
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Pred.
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                                                                                                                                                                                                                          update)
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                                                                                                                                                                                                                                                                                              86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         58
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Matches 8
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P04606;
13-AUG-1987
13-AUG-1987
16-OCT-2001
   STRAIN-ISOLATE HXB3;
MEDLINE-85228248; PubMed-2988795;
CTOW1 R., Ganguly K., Gordon M., Conroy R.,
Shaw G.M., Wong-Staal F., Reddy E.P.;
"HTLY-III env gene products synthesized in E
antibodies present in the sera of AIDS patie
Cell 41:979-986(1985).
-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT
                                                                                                                                                                                                                                                                                                                                                                                                                                           Ratner L., Haseltine W., Patarca R., Livak K.J., Starcich B.R., Josephs S.F., Doran E.R., Rafalski J.A., Whitehorn E.A., Baumeister K., Ivanoff L., Petteway S.R. Jr., Pearson M.L., Lautenberger J.A., Papas T.S., Ghrayeb J., Chang N.T., Gallo R., Wong-Staal F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus type 1 (BH10 isolate)
Human immunodeficiency virus type 1 (HXB3 isolate)
Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID-11678, 11707;
                                                                                                                                                                                                                                                                                                                                                                      "Complete nucleotide sequence of the AIDS virus, HTLV-III."; Nature 313:277-284(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=ISOLATE BH10;
MEDLINE=85111123; PubMed=2578615;
                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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InterPro; IPRO01831; HIV_Tat.
Pfam; PF00539; Tat; 1.
PRINTS; PR00055; HIVTATDOMAIN.
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FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation updat
(Transactivating regulatory pro
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Pred. No. 0.07
3; Mismatches
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patients.";
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TAT_HV1BR
P04610;
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13-AUG-1987
16-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                     Buckler C.E., Buckler-White A.J., Willey R.L., McCoy J.;
Submitted (JUN-1988) to the EMBL/GenBank/DBJ databases.
-i-FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.

MEDLINE-85099333; PubMed-2981635;
Wain-Hobson S., Sonigo P., Danos O., Cole S., Alizon
"Nucleotide sequence of the AIDS virus, LAV.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A. (CLONE PNL4-3).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Viruses; Retroid
NCBI_TaxID=11686;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cell 40:9-17(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus type 1 (BRU isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAT protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV; M15654; TAT$BH102.
HIV; M14100; TAT$HXB3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                      ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                                                                                                                                                                                                                                                                                                                      PROMOTER.
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
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Pred. No. 0.074;
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DT 15-OU
DT TAT F
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P04608; O09778;
13-AUG-1987 (Rel
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SEQUENCE
                                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rather L., Fisher A., Jagodzinski L.L., Mitsuya H., Liou R.-S., Gallo R.C., Wong-Staal F., Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.
-!- EUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE L1
                                                                                                                                                                                                     EMBL; K03455; AAB50256.1;
                                                                                                                                                                                                                                                                 or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SUBUNIT: BINDS CYCLLN TI (BY SIMILARTTY).
-!- SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ratner L., Fisher A., Jagodzinski L.L., Mitsuya H.,
Gallo R.C., Wong-Staal F.;
"Complete nucleotide sequences of functional clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1 (HXB2 isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus. NCBI_TaxID=11706;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1. PRINTS; PR00055; HIVTATDOMAIN
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HIV; M19921; TAT$NL43.
                                                                                                  InterPro;
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15-JUL-1999 (Rel. 38, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                         AF033819; AAC82591.1; K03455; TAT$HXB2.
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                                                           Pro; IPR001831;
PF00539; Tat; 1
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T -> M (IN CLONE PNL4-3).

PPOG -> AHON (IN CLONE PNL4-3).

V -> A (IN CLONE PNL4-3).

P -> S (IN CLONE PNL4-3).

9B1B4A915FAF8A14 CRC64;
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0.074;
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Pfam;

Transcription

regulation; Activator; RNA-binding; Nuclear protein;

PRINTS; PR00055; HIVTATDOMAIN

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RESULT 8
TAT_HV12%
ID TAT_H
AC P1250
DT 01-OC
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Best Local Similarity
Watches 8; Conserv
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Best Local Similarity
Matches 8; Conserv
TAT_HV1Z2
P12506;
01-OCT-1989
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P04607;
13-AUG-1987
13-AUG-1987
16-OCT-2001
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Nucleic acid structure and expression of the human AIDS/lymphadenopathy retrovirus.";
Nature 313:450-458(1985).
-1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1. PRINTS; PR00055; HIVTATDOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; K02083; AAB59870.1; -.
EMBL; X01762; -; NOT_ANNOTATED_CDS
HIV; K02083; TAT$PV22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A. MEDLINE-85111157; PubMed-2982104; Muesing M.A., Smith D.H., Cabradilla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Viruses; Retroid
NCBI_TaxID=11700;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus type 1 (PV22 isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
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Pred. No.
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Best Local
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P04609;
13-AUG-1987
13-AUG-1987
16-OCT-2001
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Submitted (NOV-1988) to the HIV data bank.

-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM TH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1 (Z2/CDC-Z34 isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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16-OCT-2001 (Rel. 40, Last annotation update)
TAT protein (Transactivating regulatory protein).
                                                                                                           domains in the envelope gene.";

Gene 52:71-82(1997).

Gene 52:71-82(1997).

Gene 52:71-82(1997).

TEUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE

TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND

ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LT
                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE-87248097; PubMed-3036660;
MIDIATUS-87248097; PubMed-3036660;
MIDIATUS-87248097; PubMed-3036660;
MIDIATUS-87248097; PubMed-3036660;
MIDIATUS-87248097; MIDIATU
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"Molecular characterization of human immunodeficiency virus from
Zaire: nucleotide sequence analysis identifies conserved and variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Viruses; Retroid viruses; Retroviridae; Lentivirus NCBI_TaxID-11708;
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SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY). SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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                                                                                  PROMOTER.
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P04613;
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13-AUG-1987
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                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alizon M., Wain-Hobson S., Montagnier L., Sonigo P.;
"Genetic variability of the AIDS virus: nucleotide sequence analysis
of two isolates from African patients.";
Cell 46:63-74(1986).
-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
                                                                                                                                                                                                                                                                                                                                 entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Viruses; Retroid
NCBI_TaxID=11697;
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Pfam; PF00539; Tat; 1.
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       SEQUENCE
                                                                                                                  PRINTS;
                                                                                                                                                                                                                                                        EMBL; X04415; CAA28015.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=86245056; PubMed=2424612;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus type 1 (MAL isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                                                                    Transcription regulation; Activator; RNA-binding; Nuclear
                                                                                                                                                                                       InterPro;
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                                                                                                                                                                        K03456; TAT$MAL.
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
(Transactivating regulatory protein).
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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P04614;
13-AUG-1987
13-AUG-1987
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
MEDLIND-85090453; PubMed-2578227;
Sanchez-Pescador R., Power M.D., Barr
Stemplen M.M., Brown-Shimer S.L., Gee
Levy J.A., Dina D., Luciw P.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; K02007; AAB59879.1; -. HIV; K02007; TAT$SF2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Science 227:484-492(1985)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Transcription regulation; Activator; RNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1. PRINTS; PR00055; HIVTATDOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus type 1 (ARV2/SF2 isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TAT protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=11685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ARV-2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'Nucleotide sequence and expression of an
                                                                                                                                                                                                                                                             47
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                                                                                                                                                                                                                                                                                                       1 YGRKKRRQKKK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     بر
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY). SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
                                                                                                                                                                                                                                                          YGRKKRRQRRR 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      YGRKKRRQRRR
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8; Conser
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8; Conserv
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(Rel.
(Rel.
                                    (Rel. 09, Created)
(Rel. 09, Last sequence update)
(Rel. 40, Last annotation update)
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                      (Transactivating
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                                                                                                                                         STANDARD;
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40,
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72.7%;
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72.7%;
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Last annotation updat
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                      regulatory protein).
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Pred. No. 0.
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                                                                                                                                                                                                                                                                                                                                                                            DB 1;
0.087;
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0.075;
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Renard A., Ra
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RESULTING THE PROPERTY OF THE 
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P20879;

01-FEB-1991

01-FEB-1991

16-OCT-2001
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Best Local
                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.

KOYANA91 S., Chen I.S.Y.;

Submitted (DEC-1988) to the HIV data bank.

-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Molecular cloning and primary nucleotide sequence analysis of a distinct human immunodeficiency virus isolate reveal significant divergence in its genomic sequences.";

Proc. Natl. Acad. Sci. U.S.A. 83:8380-8384(1986).

-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO TRANS-CTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=87041461; PubMed=3490666;
Desai S.M., Kalyanaraman V.S., Casey
Andersen P.R., Devare S.G.;
                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Viruses; Retroid viruses; NCBI_TaxID-11688;
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus type 1 (JRCSF isolate) (HIV-1).
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Pfam; PF00539; Tat; 1.
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7 YGRKKRRQRRR 57
                                                                                                                                                          SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY). SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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M13137; TAT$CDC45.
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8; Conser
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(Rel. 17, Last sequence update)
(Rel. 40, Last annotation update)
(Transactivating regulatory prote
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Bioinformatics
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Pred. No.
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Best Local S
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Gurgo C., Guo H.-G., Franchini G., Aldovini A., Collalti E.,
Farrell K., Wong-Staal F., Gallo R.C., Reitz M.S. Jr.;
"Envelope sequences of two new United States HIV-1 isolates.";
"Involopy 164:531-536(1988).
1-F FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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HIV; M38429; TAT$JRCSF.
InterPro; IPR001331; HIV_Tat.
Pfam; PF00539; Tat; 1.
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              AIDS
                                                   InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1.
                                                                               EMBL; M17449; AAA448
HIV; M17449; TAT$MN.
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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Transcription regulation; Activator; RNA-binding; Nuclear protein;
SEQUENCE
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                                         PRINTS; PR00055; HIVTATDOMAIN
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SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
MISCELLANEOUS: THE MN ISOLATE WAS TAKEN F
                                                                                                                                                                                                                                 PATIENT IN 1984.
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MEDLIND=90148544; PubMed=2559749;

Huet T., Dazza M.C., Brun-Vezinet F., Roelants G.E., Wain-Hobson S.;

"A highly defective HIV-1 strain isolated from a healthy Gabonese individual presenting an atypical western blot.";

AIDS 3:707-715(1989).

-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LITR
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-!- SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
-!- MISCELLANEOUS: THE OYI ISOLATE WAS TAKEN FROM THE BLOOD OF A
HEALTHY GABONESE INDIVIDUAL.
-!- MISCELLANEOUS: THE SINGLE C -> S SUBSTITUTION AT RESIDUE 22 OF THE
OYI TAT PROTEIN RENDERS IT INACTIVE, BUT MAY NOT ACCOUNT FOR THE
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HIV; M26727; TAT$OYI.
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MEDLINE-1322025; pubMed=11429117;

MEDLINE-1322025; pubMed=11429117;

MEDLINE-1322025; pubMed=114 V.R.K., Ahmad N.;

"Characterization of HIV Type 1 tat Sequences Associated with Perinatal Transmission ";

Perinatal Transmission";

AIDS Res. Hum. Retroviruses 17:765-773(2001).

-i-FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).

EMBL; AV007056; AAG32103.1;

-InterPro: IPR001831; HIVTATIONAIN.

PRINTS; PR00539; Tat; 1.

PRINTS; PR00539; Tat; 1.
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NON_TER 72 72
SEQUENCE 72 AA; 8342 MW;
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1 YGRKKRRQKKK 11
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9; Conserv
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(TRANSACTIVATING REGULATORY PROTEIN) (F
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MEDLINE-21322025; Pubmed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
"Characterization of HIV Type 1 tat S
Perinatal Transmission.";
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EMBL; AY007058; AAG32105.1; -
Interpro; IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1.
PRINTS; PR00055; HIVTATOMAIN.
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-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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                                   PROMOTER (BY SIMILARITY).
EMBL; AY007059; AAG32106.1; -.
In\underpro; IPR001831; HIV_Tat.
Pfam; PP00539; Tat; 1.
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MEDLINE-21322025; PubMed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
"Characterization of HIV Type 1 tat So
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
       PRINTS;
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FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
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(TrEMBLrel. 19, 1
(TRANSACTIVATING
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(TrEMBLrel. 19, Last annotation update)
(TRANSACTIVATING REGULATORY PROTEIN)
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ATING REGULATORY PROTEIN) (FRAGMENT).
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Pred. No. 0.
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PROTEIN) (FRAGMENT).
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NON_TER
SEQUENCE
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SEQUENCE FROM N.A.
STRAIN-PAIRD, MOTHER;
MEDIANE-21322025; PubMed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
                                                                                                                                                  Human immunodeficiency virus type 1. Viruses; Retroid viruses; Retrovirio NCBI_TaxID=11676;
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Pfam; PF00539; Tat; 1.
PRINTS; PR00055; HIVTATDOMAIN.
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AIDS Res. Hum. Retrovirus
-!- FUNCTION: TRANSCRIPTI
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MEDLINE-21322025; PubMed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
"Characterization of HIV Type 1 tat S
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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S RES. Hum. Retrovituses 17:765-773(2001).

FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM PROMOTER (BY SIMILARITY).

L; AY007060; AAG32107.1; -.
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1 (TrEMBLrel. 17, La
1 (TrEMBLrel. 19, La
1 (TRANSACTIVATING)
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(TremBlrel. 17, Last sequence update)
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G REGULATORY PROTEIN) (FRAGMENT).
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2; Mismatches
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EMBL; AY007063; AAG32110.1; -.

InterPro; IPR001831; HIV_Tat.

Pfam; PF00539; Tat; 1.

PRINTS; PR00055; HIVTATDOMAIN.
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AIDS Res. Hum. Retroviruses 17:765-773(2001).
1-1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-PAIRD, MOTHER;
MEDLINE-21322025; PubMed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
Characterization of HIV Type 1 tat Se
                                                                              Q99AN9
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AIDS Res. Hum. Retroviruses 17:765-773(2001).
-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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NCBI_TaxID=11676;
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EMBL; AY007062; AAG32109.1; -
InterPro; IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1.
PRINTS; PR00055; HIVTATDOMAIN.
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                                                                           PRELIMINARY;
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REGULATORY PROTEIN) (FRAGMENT).
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Pred. No. 0.
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Best Local :
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STRAIN-PAIRD, MOTHER;

MEDILINE-21322025; PubMed-11429117;

HUSAIN M., Hahn T., Yedavalli V.R.K., Ahmad N.;

"Characterization of HIV Type 1 tat Sequences Associated with Perinatal Transmission.";

ALDS Res. Hum. Retroviruses 17:765-773(2001).

-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).

EMBL; AY007067; AAG32114.1; -.

TRANSCRIPTION THE TRANSCRIPTION THE LTR
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Q99AN8;
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EMBL; AY007066; AAG322113.1; -.
InterPro: IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1.
PRINTS; PR0055; HIVTATIOMAIN.
                                                                                                                                                          Activator; Nuclear protein; NON_TER 72 72 72 SEQUENCE 72 AA; 8352 MW;
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PRINTS; PR00055; HIVTATDOMAIN.
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Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11676;
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STRAIN=PAIRD, MOTHER;
MEDLINE=21322025; PubMed-11429117;
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TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN) (FRAGMENT).
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                                    MEDILINE-95032920; PubMed-7946099;
Sabino E., Pan L.Z., Cheng-Mayer C., Mayer A.;
Sabino E., Pan L.Z., Cheng-Mayer C., Mayer A.;
"Comparison of in vivo plasma and peripheral blood mononuclear cell HIV-1 quasi-species to short-term tissue culture isolates: an analys of tat and C2-V3 env regions.";
AIDS 8:901-909(1994).
-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
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MEDLINE-21322025; PubMed-11429117;
HUSAIN M., Hahn T., Yedavalli V.R.K., Ahmad N.
"Characterization of HIV Type 1 tat Sequences
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Viruses; Retroid viruses; Retroviridae;
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FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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SEQUENCE
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Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: TRANSCRIPTIONAL RECULATOR THAT ACTS BY BINDING TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AN ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM ACTIVATES TRANSCRIPTION INITIATION AND/OR
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Human immunodeficiency virus type 1. Viruses; Retroid viruses; Retroviridae; NCBI_TaxID=11676;

Lentivirus

SEQUENCE FROM STRAIN-T10;

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Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RIAN ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).

EMBL; U57282; AABL7842.1; -.
InterPro; IPR001821. """-"
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01-MAY-1997
01-DEC-2001
     Q9YP33
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"LTR and tat variability of HIV-1 isolates from patients with divergent rates of disease progression.;
Virus Res. 57:11-20(1998).
-1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).
EMBL: U80540: AACB2979.1; -
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STRAIN=PATIENT LTS1;
MEDLINE-99049627; PubMed-9833881;
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Pfam; PF00539; Tat; 1.
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YGRKKRRQRRK
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(TremBLrel. 03, Last sequence update)
(TremBLrel. 19, Last annotation update)
(TRANSACTIVATING REGULATORY PROTEIN) (FRAGMENT).
     PRELIMINARY;
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81.8%;
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01-MAY-1999
01-MAY-1999
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                                                                                                                                                                             STRAIN-ARMA185;
                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                          Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11676;
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN).
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Burger H.;
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19 AAW76156		LB AAW31210		AAR4418	AAB7175	AAB7175				AAB7330	AAB6767	AAG7045	AAG6837			22 AAE03730			22 AAE05268			22 AAB82757	AAB60	AAE12	AAE1260	AAME	AAG6567			AAU0981	AAU0993	AAY71	AAY9354
TAT	of the i		Residues 47-58 HIV	Anti-herpetic pept	Ä	NTR3 derived pepti	protein	protein	t pr	HIV-1 TAT protein	duct		₽	transduc			Human immunodefici	HIV TAT peptide fr		Human Immunodefici	t protei	HIV TAT protein tr	Internalising pept	Membrane transport	numi	-1 tat protei	at protein f	immunodefic	in tr	HIV-1 tat protein	Human immunodefici	immun	Amino acid sequenc

ALIGNMENTS

RESULT AAU09906

AAU09906 standard; Protein;

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Jing WPI; 2001-611392/70. 16-MAR-2000; 28-NOV-2000; WO200168859-A2 Synthetic. anti-convulsant; anti-asthmatic; dermatological; renal; osteopathic; vascular; cytostatic; anti-leukaemic; anti-infertility; ophthalmological; hepatitis; anorexia; cachexia; neuronal dysfunction; lung disease; (AMGE-) AMGEN INC. 15-MAR-2001; 2001WO-US08678. 20-SEP-2001. bone s disease; 2000US-189816P. 2000US-0724460. vascular disorder; eye disorder; cancer; human.

Interleukin 17; hIL-17 receptor like protein; immunomodulatory; anti-inflammatory; anti-diabetic; immunosuppressive; anti-microbial; hepatic; anabolic; anorectic; anti-lazheimer's; anti-parkinsonian; anti-convulsant; anti-asthmatic; dermatological; renal; osteopathic;

Interleukin 17 (hIL-17) receptor like protein associated peptide #1

14-FEB-2002 AAU09906;

(first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
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Note: This artificial peptide sequence is given in the sequence listing but is not described in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acids encoding interleukin 17 receptor like polypeptides, useful for preventing, diagnosing and treating, e.g. leukemia, asthma diabetes, psoriasis and glaucoma –
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 149; 158pp; English
                                                                                                                                                                                                                                                                                                                      Sequence deduced from tat gene of HIV 1-NDK.
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR09303 standard; protein; 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor like polypeptides useful as vaccines and in gene therapy. These have immunomodulatory, anti-inflammatory, anti-diabetic, anti-microbial,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention describes novel nucleic acids encoding interleukin (IL) 17
02-MAY-1990;
                                         15-NOV-1990
                                                                                                                                         Region
                                                                                                                                                                               Region
                                                                                                                                                                                                                                             HIV 1-NDK.
                                                                                                                                                                                                                                                                             Human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                27-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                        AAR09303;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunosuppressive, hepatic, anabolic, anorectic, anti-alzheimer's, anti-parkinsonian, anti-convulsant, anti-asthmatic, dermatological,
                                                                                 WO9013630-A
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nes 9; Conserv
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ygrkkrrqrrk 11
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    90WO-FR00312
                                                                                                                                                          /label - tat ex2
                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                         /label= tat ex3
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Pred. No.
                                                                                                                                                                                                                                                                                   AIDS
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The HIV NDK virus was isolated from peripheral blood lymphocytes of an AIDS patient. A genomic library was prepd. from DNA extracted from CEM cells infected with the virus. The library was screened with a pBTl probe corresp. to a fragment from HIV 1. The virus is more cytopathic than other strains and is not inhibited by OKT4A. It has been deposited as CNCM I-857. The sequence can be used to express proteins useful for diagnosing the presence of NDK and related viruses and in vaccines against immunodeficiency diseases. See also AAR09301-4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New HIV-NDK retrovirus and protein component - used in against immuno-deficiency disorders and in raising MAbs retro-virus detection in vivo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Barre-Sinnoussi F,
Spire B;
                                                                                                                                                                                                                                                      Mouse; BH3 interacting domain death agonist; BID; BCL-2 family; apoptosis; regulation; cell death; inflammation; cancer; arthriautoimmune disease; viral infection; lymphoproliferative.
                                                                                                                                                                                                                                                                                                                                                       AAW50263;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 2; 37pp; French
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-MAY-1989;
                     BH3 interacting domain death agonist polypeptide - used for treating decreased apoptotic conditions resulting from inflammation etc.
                                                                                  Korsmeyer
                                                                                                                                 09-SEP-1996;
                                                                                                                                                         09-SEP-1997;
                                                                                                                                                                                                        W09809980-A1
                                                                                                                                                                                                                                                                                                     HIV-1 tat protein.
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                                                                                                                                                                                                                                                                                                                                                                               AAW50263 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                          WPI; 1998-193546/17
                                                                                                         (UNIW ) UNIV WASHINGTON
                                                                                                                                                                                 12-MAR-1998
                                                                                                                                                                                                                               Human immunodeficiency virus type
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                                                                                                                                  96US-0706741
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                                                                                                                                                                                                                                                                                                                                                                               Protein;
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Pred. No. 0.
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MAbs for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sire
                                                                                                                                                                                                                                                                    arthritis;
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Example 8;

Page 85; 118pp; English

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RESULT
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AC AAY2
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AC AAY2
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Best Local :
This invention describes a novel anti-pathogen system (APS) comprising a fusion protein constructed from a covalently linked protein transduction domain (PTD) and a cytotoxic domain. The APS can be used for suppressing a pathogen infection in a mammal. The method may further comprise
                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-pathogen; fusion protein; protein transduction domain; PTD; AZT; cytotoxic domain; suppressor; infection; medicament; ddI; ddC; d4T; 3TC; FTC; DAPD; 1592U89; C592; acyclovvir; ganciclovvir; peniclovir; interferon; apoptosis; virus; HIV; cytomegalovirus; CMV; herpes simplex virus; HSV-1; hepatitis virus; Kaposi's sarcoma-associated herpes virus; KSHV; berpes virus; yellow fever virus; flavivirus; rhinovirus; plasmodial;
                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dowdy
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10-DEC-1997;
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                                                                                                                                                                                                                                                                    New anti-pathogen systems, infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transduction efficiency; cytotoxin.
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                                                                                                                                                                                           34; 123pp; English
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97US-0069012
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72.7%;
                                                                                                                                                                                                                                                                                                      particularly for virus
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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RESULT
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This sequence represents a Tat peptide. The invention relates to a bcl homology domain 3 (BH3 domain), derived from a proapoptotic member of the BCL-2 family. The BH3 polypeptide can be used in a method for promoting apoptosis target cell, especially where the cell is a cancer cell a virus cell or an autoantibody producing cell. The BH3 polypeptide can
                                                                                                                                                                                                                                                                                                                                                                                                            07-OCT-1997;
26-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BH3 domain; cell death agonist; bc1 homology domain; BCL-2 family; apoptosis promoter; cancer cell; virus infected cell; inflammation; autoantibody producing cell; cancer; lymphoproliferative condition;
                                                                                                                                                                  Claim 7; Page 67; 104pp; English.
                                                                                                                                                                                                                 Bcl homology domain 3 polypeptide
                                                                                                                                                                                                                                                                  WPI; 1999-255058/21
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97US-0060133.
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                                                   beta-catenin are described. The modulating agent comprises an internalisation molety and one or more of an amino acid sequence SYLDS(PC_4)GIHS(PC_4)G, or a peptide analogue or peptidomimetic of the amino acid sequence. The modulating agents are useful for the manufacture of a medicament for stimulating hair growth or reducing hair loss, stimulating skin exfoliation, and inhibiting the development of Alzheimer's disease. They may also be used to increase the beta-catenin level in a cell, to stimulate activation of gene transcription in a cell, and to stimulate cell differentiation. They may further be used to ameliorate hearing loss resulting from a variety of inner ear disorders, such as hyperacusis and tinnitus, through regeneration of hair cells of the inner ear.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modulating agent; beta-catenin; hair loss; hair growth; skin; exfoliation; Alzheimer's disease; gene transcription; cell differentiation; hearing loss; inner ear; hyperacusis; t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in therapeutic compositions for treating disease including cancer, other lymphoproliferative conditions, arthritis, inflammation, and autoimmune diseases, which may result from the down regulation of cell death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modulating agents for inhibiting degradation of cytoplasmic beta-catenin, used for e.g. stimulating hair growth or reducing loss, inhibiting development of Alzheimer's disease, comprise internalization moiety and amino acid sequence
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                                                                                                   CC A chimeric group or fusion peptide which comprises a portion of an CC anti-apoptotic polypeptide which inhibits apoptosis of lymphocytes in CC combination with a transport group is described. The transport group is capable of transporting the chimeric group or fusion peptide is capable of transporting the chimeric group or fusion peptide is FLICE-like CC coross the cell membrane. The anti-apoptotic polypeptide is FLICE-like CC inhibitor protein (FLIP) which inhibits Fas and TNF mediated apoptosis CC by inhibiting binding of Caspase-8 to the Fas receptor complex, thus CC shutting off the downstream Fas signalling pathway. The chimeric group CC and fusion peptide are useful for inhibiting ligand-induced apoptosis CC by bringing them into contact with T cells. The chimeric group is CC useful for expanding T cells in vitro e.g. T cells specific for CC particular antigens such as tumour-specific antigen for enhancing CC immune response and to inhibit the apoptosis of chronically activated CT cells e.g. activated CD4^+ T cells in HIV infected patients. The CC chimeric group is also useful for therapeutic, prophylactic or CC diagnosis of intracellular delivery of small molecules and mucleic acids encoding such polypeptides. Two primers (AAA54298) were used to amplify the FLIP cDNA for subcloning into the Xhol-Ncol CC site of the pHA-TAT vector which contains the N-terminal protein CC site of the pHA-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CC site of the pha-TAT vector which contains the N-terminal protein CCC site of the pha-TAT vector whi
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06-APR-1999;
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                                                                                                                                                                                                                                                                                   The present invention is concerned with methods of modulating the amount of free beta-catenin in the cell, and methods of stimulating the expression of genes involved in cellular differentiation, the transcription of which is under the control of beta-catenin. The peptides given in AAB27053-B27088, AAB27284-B27300 and AAB27330-B27351 can be used as modulating agents which interrupt the interaction between alpha and beta catenin, causing increased levels of the latter and stimulating the activation of beta-catenin mediated transcription. This can be used to stimulate cell differentiation, which can then be used to promote hair growth and skin exfoliation. This latter is particularly useful in the improvement of photodamaged skin and to minimise wrinkles. The modulating peptide can also be used to reduce hearing loss resulting from inner ear disorders such as hyperacusis and tinnitus.
                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stimulating beta-catenin mediated gene expression, cellular differentiation and hair growth, involves contacting cells with modulating agent capable of inhibiting interaction between alpha beta catenin
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Protein transduction domain; fusion molecule; therapeutic agent; drug drug targetting; drug discovery; cell transduction; bioavailability; vaccine; nervous system discover; latheimer's disease; Parkinson's disease; Huntington's disease; pre-senile dementia; epiloseizure; compulsive behaviour; meningitis; encephalitis; ischaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for modulating beta-catenin mediated gene transcription in a cell. The method involves contacting a cell with a modulating agent comprising an internalization molety and a peptide comprising a sequence LXXLL or peptide analogue of LXXLL. The method is useful for modulating beta-catenin mediated gene transcription, cell differentiation, hair growth, and retinoic acid activity, for treating cancer, and for inhibiting the development of Alzheimer's disease.
                                                                                                                                                                                       HIV TAT
                                                                                                                                                                                                                                           09-FEB-2001
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                                                                                                                                                                                                                                                                                                       AAB29413;
                                                                                                                                                                                                                                                                                                                                                          AAB29413 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200063246-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                 transduction
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0296089
                                                                                                                                                                                                                                                                                                                                                          peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84.5%;
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                                                                                                                                                                                    domain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 49; DB 21 Pred. No. 0.12; 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                          11
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                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                                                                                                                          B
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                                                                                                                                                                                 ID NO:2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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RESULT , 11
AAB09907
ID AAB099
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                                                                                                                                                                                                                                                                                       Created and setting an allower to a movel fusion molecule comprising at least cone protein transduction domain (PTD) and at least one linked molecule, where the linked molecule has therapeutic or prophylactic activity against a medical condition. The invention also relates to methods of drug discovery in which the test compound is linked to a suitable createstant microorganisms using a suitable fusion molecule; a method of killing comprising a covalently linked fusion molecule; a mammal adapted for createstant microorganisms using a suitable fusion molecule; a mammal comprising a covalently linked fusion molecule; and a mammal adapted for transduced into essentially all the cells of the mammal. The fusion compound is linked molecule has been transduced into essentially all the cells of the mammal. The fusion can be compound to the mammal of the fusion of a cardiovascular drug, an antitumour drug, an anti-infective drug, a cardiovascular drug, an antitumour drug, an analyssic, an ential nervous system (2000) disorder is especially Alzheimer's disease, can tainfilammatory, a disease, thuntington's disease, and also includes pre-senile compound to the prilepsy and seizures, compoulsive behaviour, meningitis, especially Alzheimer's disease, and also includes pre-senile common to the prilepsy and seizures, compoulsive behaviour, meningitis, the methods are a used to treat or prevent bacterial or viral infections. The methods are a highly effective means for transducing a molecule into an entire mammal or into specific cells, tissues, organs and conditions of the target cell of the providing providing efficient transduction of the target cell. The present sequence conficient transduction of the target cell. The present sequence conficient transduction of the target cell. The present sequence conficient transduction of the target cell. The present sequence
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                                                                                                                                                                                                               Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fusion molecules comprising protein transduction domains therapeutic agents, useful for treating e.g. Alzheimer's Parkinson's diseases, dementia and epilepsy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dowdy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UNIW ) UNIV WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-FEB-1999;
29-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            spongiform encephalopathy; dyslexia; age-related memory loss; tou Gehring's disease; viral infection, HIV; bacterial infection.
                 AAB09907 standard; peptide;
                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                     1 YGRKKRRQKKK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000-647439/62
                                                                                                               ygrkkrrgrrr 11
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                                                                                                                                                                                         l Similarity
8; Conserv
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                                                                                                                                                                                           Conservative
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99US-0151291.
                                                                                                                                                                                                               84.5%;
72.7%;
                       片
                                                                                                                                                                                                             Score 49; I
                       ⋛
                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                               DB 21;
0.12;
                                                                                                                                                                                               0
                                                                                                                                                                                                                                 Length 11
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                                                                                                                                                                                               Gaps
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RESULT 12
AAY93542
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                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is the tat protein transduction domain fron HIV, CC which can be used as a targeting signal. It can be used to internalise CC sequences, such as MUC-1 antagonists, within the cell. MUC-1 is a CC immunosuppressor, and antagonists act to reduce overactive immune CC responses. Thus, the peptide can be used to treat inflammatory disorders CC systemic lupus erythematosus, psoriasis, allergic contact dermatitis and CC systemic lupus erythematosus, polyarteritis nodosa, Goodpastures CC systemic lupus erythematosus, polyarteritis nodosa, Goodpastures CC syndrome, isopathic thrombocytopoenic purpura, autoimmune haemolytic cancenia, Graves' disease, rheumatic fever, pernicious anaemia, CC uniquis, viral myocarditis, autoimmune thyroiditis, male infertility, CC sarcoidosis, allergic encephalomyelitis, multiple scierosis, Sjorgens CC disease, Reiter's disease, Celiac disease, sympathetic ophthalmia and CC primary biliary cirrhosis, immune disorders, graft versus host disease
                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of agent capable of intracellularly inhibiting mucin MUC-1 for inducing T-cell-based immunosuppression and for treating autoimmuno disorders, transplant rejection and inflammatory disorders \,\cdot\,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200034468-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Targeting signal; MUC-1; immunosuppression; autoimmune
immune disorder; inflammatory disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV tat protein transduction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB09907;
                                                                                                                                                        AAY93542 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 15; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Agrawal B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BIOM-) BIOMIRA INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUN-2000
cytotoxic domain; pathogen infection; r plasmodial infection; cancer; prostate
                               Protein transduction system; protein transduction domain;
                                                            Amino acid
                                                                                          25-SEP-2000
                                                                                                                         AAY93542;
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1 ygrkkrrgrrr 11
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                                                                                                                                                                                                                                                                                                                                                                                                          transplant rejection.
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                                                                                                                                                                                                                                                                                                Similarity
8; Conser
                                                           sequence of a synthetic protein transduction domain.
                                                                                                                                                                                                                                                                                                                                                                              11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Longenecker
                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                               84.5%;
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                                                                                                                                                                                                                                                                                                               Score 49; I
Pred. No. 0.
                retroviral infection;
                                                                                                                                                                                                                                                                                                                                DB 21;
                                                                                                                                                                                                                                                                                                                   . 12;
                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                Length 11;
                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             autoimmune
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                                                                                                                                                                                                                                                                                                    Gaps
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cancer

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RESULT
AAY7101
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                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY93542-51 represent synthetic protein transduction domains, which are used in the protein transduction system of the invention. The specification describes a protein transduction system, which comprises a fusion protein. This fusion protein has a covalently linked protein transduction domain and cytocoxic domain. The system is useful for treating pathogen infection in mammals, infections such as those caused by CMV, HSV-1, HCV, KSHV, yellow fever virus, flavivirus or rhinovirus, retroviral infections such as HTV-1, HTV-2, HTVL-3 and/or LAV, plasmodial infections associated with P.faciparum, P.vivax, P.ovale, P.malariae. It is also useful for treating cancer, especially prostate cancer.
                                                                                                                                                       PROSPHOLAMBAN; PLB; sarcoplasmic reticulum Ca 2+ ATPase; SERCA2a; HIV TAT protein; penetratin; transport peptide; cardiomyocyte; inhibitor; cargo peptide; contractilin; cardiac contractility; cardiant; treatme cardiac disease; heart failure; muncardial disease;
           02-NOV-1998;
27-JUL-1999;
                                                                                                     WO200025804 - A2
                                                                                                                              Human immunodeficiency virus
                                                                                                                                                                                                                                                                              AAY71015;
                                                                                                                                                                                                                                                                                                    AAY71015 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein transduction system for treating cancer and pathogenic infections has a fusion protein comprising a protein transduct: domain covalently linked to a cytotoxic domain .
                                                 02-NOV-1999;
                                                                          11-MAY-2000
                                                                                                                                                                                                                     Human immunodeficiency virus TAT peptide
                                                                                                                                                                                                                                                  29-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UNIW ) UNIV WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                    1 YGRKKRRQKKK 11
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1 Ygrkkrrqrrr 11
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                                                                                                                                                                                                                                                                                                                                                                                                                        h 84.5%;
Similarity 72.7%;
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11
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                                                                                                                                                                                                                                                 (first entry)
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           98US-0106718
99US-0145883
                                                99WO-US25692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US29289.
                                                                                                                                                                                                                                                                                                                                                                                                                        Score 49; DB
Pred. No. 0.12
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transduction
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                                                                                                                                                                  treatment,
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 14
AAU09932
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cc small peptide complexes and recombinant proteins, that induces compospholamban (PLB) deficiency and inhibits the interaction between PLB cand sarcoplasmic reticulum Ca 2+ AFPase (SERCA2a) within cardiomyocytes. CT The peptide complex comprises of transport peptide like penetratin and cargo peptide selected from mutant PLB, native PLB or antibody against PLB protein (contractilin), Penetratin-PLB peptide functions as a contractility and reduces blood pressure. This method is useful for the treatment of cardiac disease e.g. heart failure and myocardial cysfunction. The present sequence is the amino terminal, ll-amino acid protein transduction domain of the denatured human immunodeficiency withus (HTV) TAT protein. It belongs to the penetratin class of peptides, compounds across the plasma membrane. It is used in the construction of CC compounds across the plasma membrane. It is used in the construction of CC recombinant PLB peptides (e.g., TAT-PLB).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                     protein transduction domain; PDT: homan '---
                         WPI; 2001-596910/67
                                                   Jing S,
                                                                                                        13-MAR-2000; 2000US-188786P
                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                   AAU09932 standard;
                                                                              (AMGE-)
                                                                                                                                   13-MAR-2001; 2001WO-US08013
                                                                                                                                                              20-SEP-2001.
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                                                                                                                                                                                                                                                                                                               15-JAN-2002
                                                                                                                                                                                                                                                                                                                                            AAU09932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating cardiac diseases, e.g. heart failure or myocardial dysfunction comprises enhancing cardiac contractility by inhibiting interaction between phospholamban and sarcoplasmic reticulum calcium 2+ adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 5; Page 54; 56pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     triphosphatase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-365393/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ygrkkrrqrrr 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 YGRKKRRQKKK 11
                                                                                                                                                                                                                immunodeficiency virus type
                                                                                                                                                                                                                                                                                immunodeficiency virus (HTV) tat partial peptide sequence
                                                                               AMGEN
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                                                    Bass MB
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Wang Y, Silverman GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            discloses a method for the treatment of heart failure, using
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                                                                                                                                                                                                                                                                                                                                                                    Peptide;
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72.7%;
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Pred. No. 0.12
3; Mismatches
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New fibroblast growth factor-like polypeptide and polynucleotide for

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RESULT 15
AAU09812
ID AAU0981
XX AAU098
XX AAU098
XX HIV-1
XX flbrob
KW flbrob
KW flbroc
KW Osteog
KW inmunc
KW diabet
KW human
XX HUMBAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fibroblast growth factor receptor-like protein; FGFR-L; anorectic; haemostatic; osteopathic; cytostatic; nephrotropic; antidiabetic; immunomodulator; antiinflammatory; haematpooletic disorder; osteoporosis; osteogenesis imperfecta; paget's disease; periodontal disease; cancer; hypercalcaemia; acute glomerulonephritis; chronic glomerulonephritis; diabetes; obesity; cachexia; transgenic animal; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis, prevention and treatment of diseases, disorders or conditions involving the central nervous system, teeth, heart, liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
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     Saris
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                                                                                                                    (XIAM/)
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                                                          BOONE T C.
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SARIS C M.
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Pred. No.
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Novel nucleic acid encoding fibroblast growth factor receptor-like polypeptides, useful for treating hematopoietic disorder, osteoporo Paget's disease, glomerulonephritis, cancer, diabetes, obesity and
                                                                                                                                                                    cachexia
                                                                                                                                                                                                 2001-626128/72.
                                                                                                                                                                                 osteoporosis,
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Disclosure; Page 59; 163pp; English.

CC hypercalcaemia, acute glomerulonephritis, chronic glomerulonephritis, cancer, diabetes, obesity and cachexia. (1) is also useful for clidentifying a compound which binds to FGFR-L polypeptide, by contacting CC (I) with a compound, determining the extent of binding of the FGFR-L polypeptide to the compound, and determining the activity of the CC polypeptide when bound to the compound. (II) is useful for modulating CC levels of a polypeptide in an animal. A transgenic animal comprising (I) cc is useful for determining whether a compound inhibits FGFR-L polypeptide CC activity or FGFR-L polypeptide production, by exposing the transgenic canimal to the compound and measuring FGFR-L polypeptide or production in CC mammal. (II) is useful for mapping the locations of FGFR-L gene and CC related genes on chromosomes, as hybridisation probes in diagnostic CC assays to test for the presence of an FGFR-L nucleic acid molecule in CC mammalian tissue or bodily fluid samples, in gene therapy, and as tools CC immunogen, and for cloning FGFR-L polypeptide genes. (I) is useful as immunogen, and for cloning FGFR-L polypeptide ligands using an CC expression cloning strategy. The present sequence represents the cc amino acid sequence of human immunodeficiency virus (HIV-1) tat protein crossion cloning companies and described in the invention. encoding (I) are useful for treating, preventing or ameliorati a medical condition including haematopoietic disorder, osteopo osteogenesis imperfecta, Paget's disease, periodontal disease, The invention relates to a novel isolated fibroblast growth factor receptor-like (FGFR-L) polypeptide (I). (I) and the nucleic acid (II) encoding (I) are useful for treating, preventing or ameliorating osteoporosis,

Sequence 11 Ä

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Query Match
Best Local S
Matches
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nes 8; Conserv
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ygrkkrrqrrr
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Search completed: July 15, Job time: 415 sec 2002, 12:57:59 밁 Š

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Result
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Maximum DB
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Perfect score:
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Listing first 45 summaries
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58
1 YGRKKRRQKKK 11
        July 15, 2002, 12:59:19; Search time 75.67 Seconds (without alignments) 3.551 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000
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        DВ
          US-08-706-741B-54
US-08-924-659A-54
US-09-296-089-37
US-09-296-089-37
US-08-450-246-47
US-08-450-236-47
US-08-450-236-47
US-08-450-236-73-47
US-08-235-403-47
US-08-235-6077-12
US-08-235-6077-11
PCT-US95-06077-12
US-08-450-257-6
US-08-450-257-6
US-08-450-257-6
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54, Appl
2, Appl
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Qy 1 Db 1	Query Matc Best Local Matches	RESULT 1 US-08-706-741B Sequence 54, Patent No. 5: Patent		22 22 23 23 23 23 23 24 24 24 24 24 24 24 24 24 24 24 24 24
YGRKKRROKKK 11 ::: YGRKKRRORRR 11	tch al Similarity 8; Conserva	741B-54 54, APPLICATION OF INVENTION OF INVE		4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4
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US-08-924-695A-54

Patent No.

Sequence 54, Application US/08924695A Patent No. 5998583

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/924,695A FILING DATE: 09-SEP-1997
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: HOLLAND, DONALD R.
REGISTRATION NUMBER: 971798
REFERENCE/DOCKET NUMBER: 971798
TELECOMMUNICATION INFORMATION:
TELEPHONE: (314) 727-5188
TELEPHONE: (314) 727-6092
INFORMATION FOR SEO ID NO: 54:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: Amino acids
                                                                                                                  GENERAL INFORMATION:
APPLICANT: DOWDY, SEGVEN F.
APPLICANT: DOWDY, SEGVEN F.
TITLE OF INVENTION: ANTI-PATHOGEN SYSTEM AND METHODS OF USE THEREOF
FILE REFERENCE: 48881/1742
CURRENT APPLICATION NUMBER: US/09/208,966
CURRENT FILING DATE: 1998-12-10
EARLIER APPLICATION NUMBER: 60/082,402
EARLIER FILING DATE: 1998-04-20
EARLIER APPLICATION NUMBER: 60/089,012
EARLIER FILING DATE: 1997-12-10
                                                                     NUMBER OF SEQ ID NOS: 5'
SOFTWARE: PatentIn Ver.
SEQ ID NO 2
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Best Local Similarity
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-09-208-966-2
          LENGTH: 11
TYPE: PRT
ORGANISM: human
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PATENTIA RELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy dlsk
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TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: ST. LOUIS
STATE: MISSOURI
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H3 INTERACTING DOMAIN DEATH AGONIST
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Pred. No. 0.067;
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TITLE OF INVENTION: COMPOUNDS AND METHODS FOR MODULATING TITLE OF INVENTION: BETA-CATENIN MEDIATED GENE EXPRESSION FILE REFERENCE: 100086.411
CURRENT APPLICATION NUMBER: US/09/296,089
CURRENT FILING DATE: 1999-04-21
NUMBER OF SEQ ID NOS: 37
SOFTWARE: FRSTEEQ for Windows Version 3.0
SEQ ID NO 37
LENGTH: 11
TYPE: PRT
ORGANISM: Human immunodeficiency virus
US-09-296-089-37
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Best Local S
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                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/450,257
FILING DATE: 25-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Blaschuk, Orest W. APPLICANT: Byers, Stephen APPLICANT: Gour, Barbara J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FRANKEL, Alan
APPLICANT: PABO, Carl
APPLICANT: BARSOUM, James G.
APPLICANT: FAMELL, Stephen E.
APPLICANT: PEPINSKY, R. B.
                                                                                                                                                                                                                                                          STREET: LALL
STREET: New York
CITY: New York
TTATE: New York
TTA
                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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APPLICATION NUMBER: US 08/235,403 FILING DATE: 28-APR-1994 APPLICATION NUMBER: US 07/934,375 FILING DATE: 21-AUG-1992
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o. 5652122
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                                                                                                                                                                                                                                                                                                                                                                                PEPINSKY, R. B.
VENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
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                                                                                                                                                       Release #1.0, Version #1.25
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Pred. No. 0.067;
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Pred. No. 0.067;
3; Mismatches
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US-08-450-246-47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 47,
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Best Local
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TELEX: 14-8367
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ...ACTERISTICS:
...YPE: amino acids
STRANDEDMESS: si-
TOPOLOGY:
OLECULF
(SCOULDE)
                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,403
FILING DATE: 28-APR-1994
APPLICATION NUMBER: US 07/934,375
FILING DATE: 21-AUG-1992
APPLICATION NUMBER: US 07/098,766
FILING DATE: 05-777-1997
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: PEPINSKY, R. B.
TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B170 CIP
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION: NAME: Haley Jr., James F
                                                                                                               APPLICATION NUMBER: FILING DATE: 25-MAN CLASSIFICATION: 51
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nes 8; Conserv
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                                                                                                                                                                                                                                                                                              CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 02-JAN-1991
APPLICATION NUMBER: US 0:
FILING DATE: 24-NOV-1993
 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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FILING DATE: 21-DEC-1989
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                                                                                                                                                                                                                                                                                                                                                                                               BARSOUM, James G. FAWELL, Stephen E.
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                                                                                                                               25-MAY-1995
            MBER: US 07/098,766
28-JUL-1993
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72.7%;
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PCT/US93/07833
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Pred. No. 0.072;
3; Mismatches
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 25-MAY CLASSIFICATION: 514
                                                                                                                                                                                                                                                         MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                  STREET:
       FILING DATE: 19-AU APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: B1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: Haley Jr., James F
   Local Similarity
hes 8; Conserv
                                                                                                                         TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                  NAME: Haley Jr., Jar
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 02-JA
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 0 FILING DATE: 21-DEC-1989
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                                                                                                                                                     12 amino acids
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   Conservative
                                                                                                              linear
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                                                                                            peptide
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02-JAN-1991
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                                                                                                                                                                                                                                                                   B170 CIP 2
                 Score 49; 1
Pred. No. 0
   Mismatches
                             Length 12;
 Indels
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1 YGRKKRROKKK 11 YGRKKRRQRRR 11

Gaps

0;

GENERAL INFORMATION:
APPLICANT: FRANKEL
APPLICANT: PABO, CI
APPLICANT: BARSOUM Sequence 47, Application US/08450098 Patent No. 5674980 APPLICANT: PEPINSKY, R. B. TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS: BARSOUM, James G. FAWELL, Stephen E. PEPINSKY, R. B. PABO, Carl FRANKEL, Alan

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA: New York New York 1251 Avenue of the Americas FISH & NEAVE Floppy disk

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08,
FILING DATE: 28-APR-1994
APPLICATION NUMBER: US 07,
FILING DATE: 21-AUG-1992
APPLICATION NUMBER: US 07,
FILING DATE: 28-JUL-1993
APPLICATION NUMBER: PCT/US UMBER: PCT/US93/07833 19-AUG-1993 MBER: US/08/450,098 25-MAY-1995 US 08/235,403 US 07/934,375 US 07/454,450 US 07/098,766

APPLICATION NUMBER: FILING DATE: 02-JAN

02-JAN-1

US 07/636,662 -1991

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; MOLECULE TYPE: peptide US-08-450-098-47
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Best Local Similarity
"~*^hes 8; Conserva
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TELEPHONE: (212) 596-9000
TELEFAX: (212) 596-9090
TELEX: 14-8367
INFORMATION FOR SEQ ID NO: 47:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 47, Apprint No. 5747641
                                                                           PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/235,403
APPLICATION NUMBER: US 07/934,375
APPLICATION NUMBER: US 07/934,375
FILING DATE: 21-AUG-1992
APPLICATION NUMBER: US 07/098,766
FILING DATE: 28-JUL-1993
APPLICATION NUMBER: PCT/US93/07833
FILING DATE: 19-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                      COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA PLACASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/451,233
FILING DATE: 25-MAY-1995
CLASSIFICATION: 514
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ATTORNEY/AGENT INFORMATION:
NAME: Haley Jr., James F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: PEPINSKY, R. B.
TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1251 Avenue of the Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Haley Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B170 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US UFFILING DATE: 24-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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                                APPLICATION NUMBER:
                                                  APPLICATION NUMBER: FILING DATE: 21-DE
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linear
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NUMBER:
                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                              21-DEC-1989
                02-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Carl
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                                US 07/636,662
                                                                  US 07/454,450
US 08/158,015
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Pred. No. 0.072;
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Best Local :
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TELEX: 14-8367
INFORMATION FOR SEQ ID NO:
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APPLICANT: PABO, Car
APPLICANT: BARSOUM,
APPLICANT: FAWELL, S
APPLICANT: PEPINSKY,
                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIA Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/450,236

FILING DATE: 25-MAY-1995

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
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REFERENCE/DOCKET NUMBER: B1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: BARSOUM, James G.
APPLICANT: FAWELL, Stephen E.
APPLICANT: PEPINSKY, R. B.
TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 24-NOV-1993
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                    APPLICATION NUMBER: PCT/US
APPLICATION NUMBER: US 07/
FILING DATE: 19-AUG-1993
APPLICATION NUMBER: US 07/
FILING DATE: 21-DEC-1999
APPLICATION NUMBER: US 07/
FILING DATE: 02-JAN-1991
ATTORNEY/AGENT NAME: Haley
                       APPLICATION NUMBER: US 08
APPLICATION NUMBER: US 08
APPLICATION 24-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: New York
STATE: New York
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nes 8; Conserv
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                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/235,403 FILING DATE: 28-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                  APPLICATION NUMBER: US 07/098,766 FILING DATE: 28-JUL-1993
                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/934,375 FILING DATE: 21-AUG-1992
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linear
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                   INFORMATION:
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                                                                            US 07/636,662
1-1991
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                                                       US 08/158,015
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                                   Query Match
Best Local Similarity
% 8; Conserv
                                                                                                                             ; MOLECULE TYPE: peptide US-08-733-505A-50
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US-08-733-505A-50
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Best Local
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: B170
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
TELEFAX: (212) 596-9090
TELEX: 14-8367
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                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: HOLLAND, DONALD R.
REGISTRATION NUMBER: 35,197
REFERENCE/DOCKET NUMBER: 96:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/733,505A
FILING DATE:
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: SERINE SUBSTITUTED MUTANTS OF
TITLE OF INVENTION: BCL-XL/BCL-2 ASSOCIATED CELL DEATH REGULATOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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CITY: ST. LOUIS
STATE: MISSOURI
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les 8; Conserv
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STRANDEDNESS: sin
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    1 YGRKKRRQKKK 11
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                                                   Conservative
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72.7%;
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72.7%;
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                                                Score 49; DB 2;
Pred. No. 0.072;
3; Mismatches
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Pred. No. 0.072;
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                                                                            Length 12;
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RESULT 12
PCT-US95-06077-4
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Query Match
Best Local Similarity
Thes 8; Conserva
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TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-235-403-47
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US-08-235-403-47
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                                                                                                                                                                                                                                                                                      TELEX: 14-8367
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                    NAME: Haley Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B170
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
TELEPAX: (212) 596-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 21-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: LLL-
STREET: New York
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION: NAME: Haley Jr., James F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: BARSOUM, James G. APPLICANT: FAWELL, Stephen E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: FRANKEL,
APPLICANT: PABO, Car
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 19-AUG-1993
APPLICATION NUMBER: US 07/454,450
FILING DATE: 21-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07
FILING DATE: 28-JUL-1993
APPLICATION NUMBER: PCT/1
FILING DATE: 19-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/636,662
FILING DATE: 02-JAN-1991
APPLICATION NUMBER: US 08/158,015
FILING DATE: 24-NOV-1993
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STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
1 YGRKKRRQRRR 11
                                1 YGRKKRRQKKK 11
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                                                                       Conservative
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VENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
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                                                                                      84.5%;
72.7%;
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                                                                                      Score 49; 1
Pred. No. 0
                                                                       Mismatches
                                                                                    DB 4; Length 12;
0.072;
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Best Local (
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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APPLICATION NUMBER: US 06
FILING DATE: 23-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
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OPERATING SYSTEM: PC-DOS/MS-DOS
SODTWARE: Patentin Rel-ease #1.0, Version #1.25
CURRENT APPLICATION_DATA:
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BR PC compatible
OPERAFING SYSTEM: PC-DOS/MS-DOS
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MOLECULE TYPE:
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CITY: S
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                                                                                                                 STREET: Spring House
CITY: Spring House
STATE: Pennsylvania
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ENTION: Vaccine Interdiction of Extracellular

ENTION: Transactivating Proteins of Human Immunodeficiency Virus

ENTION: and Other Chronically Infecting Viruses Employing Similar

ENTION: Intercellular Transactivating Strategies
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Transactivating Proteins of Human Immunodeficiency Virus
and Other Chronically Infecting Viruses Employing Similar
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PCT-US95-06077-11
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APPLICATION NUMBER: US 01
FILING DATE: 23-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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   SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
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TITLE OF INVENTION:
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STATE:
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                                                                                                                             NAME: Bak, Mary E. REGISTRATION NUMBER:
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les 8; Conserv
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Spring House
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Pred. No. 0.081;
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RESULT 15
PCT-US95-06077-12
Commence 12, Application PC/TUS9506077
Search completed: July 15, 2002, 12:59:20 Job time: 391 sec
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PCT-US95-06077-12
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; MOLECULE TYPE: peptide
PCT-US95-06077-11
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Best Local Similarity
Matches 8; Conserv
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TELEFAX: 215-540-9818
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
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Best Local Similarity
Matches 8; Conserv
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CLASSIFICATION:
PRIOR APPLICATION NUMBER: US 08/247,991
FILLING DATE: 23-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: IRI44PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
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MEDIOM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06077
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TITLE OF INVENTION: Vaccine Interdiction of Extracellular
TITLE OF INVENTION: Transactivating Proteins of Human Immunodeficiency Virus
TITLE OF INVENTION: and other Chronically Infecting Viruses Employing Similar
TITLE OF INVENTION: Intercellular Transactivating Strategies
NUMBER OF SEQUENCES: 38
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TYPE: amino acid
TOPOLOGY: unknown
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Pred. No. 0.081;
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Pred. No. 0.081;
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Perfect score:
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         protein search, using sw model
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length: 2000000000
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ALIGNMENTS

RESULT 1

S37533

Ig kappa chain V region (V-kappa 4) - human (fragment)

C;Species: Homo sapiens (man)

C;Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *text_change 23-Jul-1999

C;Accession: S37533

R;Klein, U; Kueppers, R; Rajewsky, K.

submitted to the EMBL Data Library, September 1993

A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the periphe A;Accession: S37533

A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 <KLE>
A;Cross-references: EMBL:226627; NID:9405708; PIDN:CAA81380.1; PID:9405709

C;Superfamily: immunoglobulin V region; immunoglobulin homology

C;Keywords: heterotetramer; immunoglobulin

the peripheral

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RESULT 2

S37530

Ig kappa chain V region (V-kappa 4) - human (fragment)

C;Species: Homo sapiens (man)

C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999

C;Accession: S37530

R;Klein, U; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993

A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the periphe A;Reference number: S37501

A;Reference number: S37501

A;Cross-references: EMBL: 226630; NID:9405702; PIDN:CAA81383.1; PID:9405703

C;Keywords: heterotetramer; immunoglobulin
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Ig Kappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: 337535
R;Klein, U; Kueppers, R; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human 1gM(+)IgD(+) cells, the major B cell subset in the periphe:
A;Reference number: S37501
A;Recession: S37535
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 <KLE>
A;Residues: 1-92 <KLE>
A;Residues: 1-92 <KLE>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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S37529
S37529
C;Bappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: S37529
R;Klein, U.: Kueppers, R.: Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM/JIDD(+) cells, the major B cell subset in the peripheral blood A;Reference number: S37501
A;Accession: S37529
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 KLEP
A;Cross-references: EMBL: Z26631; NID:g405700; PIDN:CAA81384.1; PID:g405701
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
RESULT 5
$37531

Ig kappa chain V region (V-kappa 4) - human (fragment)
C; Species: Homo sapiens (man)
C; Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *text_change 23-Jul-1999
C; Accession: $37531
R; Klein U; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A; Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peri A; Recession: $37531
A; Status: preliminary
A; Molecule type: mRNA
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Ig kappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *text_change
C;Accession: S37534
                                                                                                                                                            A;Molecule type: mRNA A;Residues: 1-92 <KLE9. A;Cross-references: EMBL:226628; NID:9405706; PIDN:CAA81381.1; PID:9405707 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-92 «KLE>
A;Cross-references: EMBL:226626; NID:9405710; PIDN:CAA81379.1; PID:9405711
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Accession: S37534
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                                                                                                                                                                                                                                                 R;Klein, U; Kueppers, R.; Rajewsky, K.

R;Klein, U; Kueppers, R.; Rajewsky, K.

submitted to the EMBL Data Library, September 1993

A;Description: Human IgM(+)IgD(+) cells, the major B

A;Reference number: S37501

A;Accession: S37532

A;Status: preliminary

A;Status: preliminary
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C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *te
C;Accession: S37532
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RESULT 9

B34153

Ig kappa chain V-IV region (Fue) - human
C.Species: Homo sapiens (man)
C.Date: 11-Apr-1990 #sequence_revision 11-Apr-1990 #text_change 21-Jan-2000
C.Accession: B34153

R.Mihaesco. E.; Ayadi, H.; Congy, N.; Gendron, M.C.; Roy, J.P.; Heyermann, H.; Frangione
J. Biol. Chem. 264, 21481-21485, 1989
A.Title: Multiple mutations in the variable region of the kappa light chains of three mc
A; Reference number: A34153; MUID:90094313
A; Accession: B34153
A; Accession: B34153
A; Accession: B34153
A; Status: preliminary
A; Molecule type: protein
A; Residues: 1-102 cMIH>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin homology
C; Keywords: immunoglobulin homology <IMM>
A49138

IgA kappa rheumatoid factor variable - human C; Species: Homo sapiens (man)
C; Date: 19-Dec-1993 *sequence_revision 18-No C; Accession: A49138
R; Gause, A.; Kuppers, R.; Mierau, R.
Clin. Exp. Immunol. 88, 430-434, 1992
A; Title: A somatically mutated V kappa IV ge A; Reference number: A49138; MUID:92298590
A; Accession: A49138
A; Status: preliminary
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C;Species: Homo sapiens (man)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 21-Jan-2000
C;Accession: PH0869
R;Manheimer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, B.
J. Exp. Med. 174, 1639-1652, 1991
A;Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idiotype A;Accession: PH0862; MUID:92078875
A;Reference number: PH0862; MUID:92078875
A;Accession: PH0862; MUID:92078875
A;Nolecule type: DNA
A;Residues: 1-101 <MAN>
A;Note: residues 28-33 were obtained from Figure 4
C;Comment: This antibody is produced by Epstein-Barr virus-transformed B cell that bears
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology
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F;95-101/Region: complementarity-determining
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F;56-62/Region:
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F;16-96/Domain: immunoglo
F;24-40/Region: complemen
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A;Reference number: S305;
A;Accession: S30523
A;Status; preliminary
A;Molecule type: mRNA
A;Residues: 1-113 <MAR>
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                                                                                A;Cross-references: EMBL:Z18325
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-113 <MAR>
                                                                                                                                                                                                              A; Reference number: A; Accession: S30520
                                                                                                                                                                                                                           submitted to the EMBL Data Library, A; Reference number: $30520
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <II
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R; Mariette, X.
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A; Residues: 1-106 <GALV
A; Residues: 1-106 <GALV
A; Cross-references: GB:S37926; NID:g298207; PIDN:AAB22366.1; PID:g298208
A; Note: sequence inconsistent with the nucleotide translation
A; Note: sequence extracted from NCBI backbone (NCBIN:106633, NCBIP:106637)
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F; 20-99/Domain: immunoglobulin homology <IMM>
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number: S30520
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                  Score 87;
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RESULT 14

SJ4002

Ig kappa chain V region - human
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Accession: SJ4002; SJ0522
C; Accession: SJ4002; SJ0522
R; Mariette, X: Tsapis, A.; Brouet, J.C.
Ritl: Nucleotidic sequence analysis of the variable domains of four human monoclonal
A; Reference number: SJ4001; MUID:93209281
A; Rocession: SJ4002
A; Status: preliminary
A, Molecule type: mRNA
A; Residues: 1-113 <MAR>
A; Cross -references: EMBL:218328
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology <IMM>
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C;Date: 02-Dec-1993 **sequence_revision 10-Nov-1995 **text_change 21-Jan-2000
C;Accession: S34003
R;Mariette, X; Tsapis, A.; Brouet, J.C.
Eur. J. Immunol. 23, 846-851, 1993
A;Title: Nucleotid: sequence analysis of the variable domains of four human A;Reference number: S34001; MUID:93209281
A;Reference number: S34001; MUID:93209281
A;Accession: S34003
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-113 cMAR>
C;Keywords: heterotetramer; Immunoglobulin
C;Keywords: heterotetramer; Immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
KAHULN

(RAPPA Chain V-IV region (Len) - human

() Species: Homo sapiens (man)

() C; Ste: 24-Apr-1984 #sequence_revision 24-Apr-1984 #text_change 01-Dec-2000

() Accession: A01903; F61458

R; Schneider, M; Hilschmann, N:

Hoppe-Seyler's Z. Physiol. Chem. 356, 507-557, 1975

Hoppe-Seyler's Z. Physiol. Chem. 356, 507-557, 1975

A; Title: Die Primaerstruktur einer monoklonalen Immunglobulin-L-Kette der Staten in St
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S34003
Ig kappa chain V region - human
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Search completed: July 15, Job time: 469 sec

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A;Molecule type: protein
A;Residues: 1-13 <BRO>
C;Comment: This is a Bence Jones protein.
C;Comment: An immunoglobulin heterotetramer subunit consists of two identical light (
c;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (
hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into
c;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
F;23-94/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Expression of a public idiotype by human monoclonal IgM directed to myelin-a A;Reference number: A61458; MUID:90039128 A;Accession: F61458
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Matches 18; Conservative 0;
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                              SGSGTDFTLTISSLQAED 29
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Pred. No. 0.00014;
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Copyright (c) 1993 - 2000 Compugen Ltd.
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01-JAN-1988 (Rel. 06, Created)
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15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain V-IV region precursor (Fragment).
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19 kappa chain V-IV region JI precursor.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
   SEQUENCE FROM N.A.
MEDLINE-86041853; PubMed-2997712;
Klobeck,H.G., Bornkamm G.W., Comb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-86041853; PubMed-2997712;
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Catarrhini; Hominidae;
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Nucleic Acids Res. 13:6515-6529(1985).
This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outset the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed.
                                                                                                                                                                                                                                           "Detection cDNA probe.
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01-APR-1988 (Rel. 07, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
19 kappa chain V-TV region B17 precursor.
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21-JUL-1986 (Rel. 01, Last sequence up
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19 kappa chain V-I region Hau.
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RESULT 7
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  KV1I_HUMAN
P01601;
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PIR; A01901; K3HUVH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A. MEDLINE-85087932; PubMed-6440122;
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13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
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RESULT 8
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DT 13-AUG
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DE IG KAP
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Best Local Similarity 94.4
Matches 17; Conservative
KV1W_HUMAN
P02431;
13-AUG-1987
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Ig kappa cha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Human immunoglobulin variable region genes kappa genes and a pseudogene.";
Nature 288:730-733(1980).
                                                                                                                                                                                                                                                                                                                                           Immunoglobulin
SIGNAL 1
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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EMBL; K01324; AAA58932.1; -.
EMBL, V00558; CAA23824.1; -.
PIR; A01881; K1HU11.
PIR; A21056; A21056.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens (Human)
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21-JUL-1986 (Rel. 01,
15-JUL-1999 (Rel. 38,
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ain V-I region Walker precursor.
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FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
BY SIMILARITY.
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Pred. No.
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RESULT 9

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"Immunoglobulin genes of the kappa light ci
lymphoid cell lines are closely related.";
                                                                                                                            Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation updating kappa chain V-III region POM.
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Interpro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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                                                                   MEDLINE-76276460; PubMed-60899;
Klapper D.G., Capra J.D.;
                                                                                                                                                    Homo sapiens (Human)
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Mammalia; Eutheria;
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COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
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COMPLEMENTARITY-DETERMINING-1.
                                                                                                                              Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE-81241357; PubMed-6788890;
Kwan S.-P., Rudikoff S., Seidman J.G., Leder P., Scharff M.D.;
Nucleic acid and protein sequences of phosphocholine-binding light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Piam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
Immunoglobulin V region.
DISULFID 23 89
                                                                                                                                                                                                                              the European Bioinformatics Institute. The use by non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                Ig kappa chain V-I region S107A. Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                              P01632;
21-JUL-1986 (Rel.
                                                                                                                                                                                        EMBL; U29423; AAC00033.1;
                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a coefficient the Swiss Institute of Bioinformatics and the EMBL
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                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                               21-JUL-1986 (Rel. 01, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                               KV1A_MOUSE
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InterPro; IPR003596; Ig_v.
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InterPro; IPR003006;
InterPro; IPR003596;
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mes 16; Conserv
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noglobulin V regi
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Rodentia;
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Pred. No.
                                                                    COMPLEMENTARITY-DETERMINING-3. FRAMEWORK-4. BY SIMILARITY.
                                                                                                   FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
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                                                                                                                  COMPLEMENTARITY - DETERMINING - 1
                                                                                                                          FRAMEWORK-1.
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                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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                                                    32008EC8E9DBE67B CRC64;
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21-JUL-1986 (Rel. 0
21-JUL-1986 (Rel. 0
15-JUL-1999 (Rel. 3
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DOMAIN
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Biochem. J. 171:337-347(1978),

-i- MISCELLANEOUS: THE MATURE CHAIN HAS 12 ADDITIONAL RESIDUES AT ITS

-i- MISCELLANEOUS: THE MATURE CIAIN HAS 12 ADDITIONAL RESIDUES AFTER THE

AMINO END, DUE TO A TRANDEM DUPLICATION OF 36 NUCLEOTIDES AFTER THE

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CODON FOR RESIDUE 36. RESIDUE 42 CORRESPONDS TO THE AMINO-TERMINAL
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Rabbitts T.H., Hamlyn P.H., Matthyssens G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Functional significance and evolutionary development of the 5'-terminal regions of immunoglobulin variable-region genes."; Cell 29:681-689(1982).
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MEDLINE-83001944; PubMed-6288267;
Kelley D.E., Colectough C., Perry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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   SEQUENCE
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InterPro; IPR003596; Ig_v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oetween
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Sequence of the full-length immunoglobulin kappa-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Smith G.P.;
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                                                                                         FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3
                                                                                                                                                                 IG KAPPA CHAIN V-V REGION MPC11. FRAMEWORK-1. COMPLEMENTARITY-DETERMINING-1.
                                                                           FRAMEWORK-4.
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COMPLEMENTARITY-DETERMINING-2.
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 B0480C87B682AC3E CRC64;
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13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
19 kappa chain V-III region VG precursor (F
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Hoppe-Seyler's Z. Physiol. Chem. 355:842-866(1974).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KV10_HUMAN
P01609;
                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; P
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InterPro; IPR003006;
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21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
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SMART; SM00406; IGV;
  SEQUENCE FROM N.A.
                                                 NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eulitz M., Hilschmann N.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=75059271; PubMed=4435756;
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Query Match
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Matches 16; Conser
                                                                                                                        Coapra J.D., Klapper D.G.;
"Complete amino acid sequence of the variable of IgM anti-gamma globulins (Lay/Pom) with shared specificities."
                                                                  Scand. J. Immunol. 5:677-684(1976).

-I- MISCELLANEOUS: THE SECOND AND THIRD HYPERVARIABLE REGIONS OF THIS CHAIN ARE IDENTICAL WITH THOSE OF THE HUMAN POM V-III KAPPA CHAIN, WITH WHICH IT SHARES CERTAIN IDIOTYPIC DETERMINANTS.
                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
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SMART; SM00406; IGv; 1.
Immunoglobulin V region; Signal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic Acids Res. 12:9229-9236(1984).
                                                                                                                                                                                MEDLINE=77038198; PubMed=824717;
                                                                                                                                                                                                                       NCBI_TaxID=9606;
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Immunoglobulin genes of different subgroups
                                                       MISCELLANEOUS:
                           GLOBULIN ACTIVITY. A01871; K1HULY.
A01871; K1HULY.
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Kratzin H., Yang C., Krusche J.U., Hilschmann N.;
"Preparative separation of the tryptic hydrolysate of a protein
high-pressure liquid chromatography. The primary structure of a
monoclonal L-chain of k-type, subgroup I (Bence-Jones Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KV1S_HUMAN STANDARD; PRT; 1
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21-JUL-1986 (Rel. 01, Created)
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15-JUL-1999 (Rel. 38, Last annotation
15 kappa chain V-I region Wes.
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1 23 FRAMEWORK-1.
24 34 COMPLEMENTARITY-DETERMINING-1.
35 49 FRAMEWORK-2.
50 56 COMPLEMENTARITY-DETERMINING-2.
57 88 FRAMEWORK-3.
69 97 COMPLEMENTARITY-DETERMINING-3.
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98 107 FRAMEWORK-4.
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COMPLEMENTARITY-DETERMINING-2.
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COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
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Search completed: July 15, 2002, 13:23:34 Job time: 1450 sec

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Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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SEQUENCE
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Clin. Immunol. Immunopathol. 87:184-192(1998).
EMBL; AF03503; AAD56269.1; -.
HSSP; P01607; 1REI.
InterPro; IPR003906; Ig_MHC.
InterPro; IPR00396; Ig_WHC.
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SMART; SM00406; IGV; 1.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID-9606;
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01-MAY-2000 (TrEMBLrel. 1:
01-DEC-2001 (TrEMBLrel. 1:
MYOSIN-REACTIVE IMMUNOGLO
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MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
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Adderson E.E., Shikhman A.R., Ward K.E., Cunningham M.W.;
Molecular analysis of polyreactive monoclonal antibodies from rheumatic carditis: human anti-N-acetylglucosamine/anti-myosin antibody V region genes.";
J. Immunol. 161:2020-2031(1998).
EMBL; U96396; AAB68785.1; -.
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Q1-DEC-2001 (TrEMBLrel. 19, Created)
Q1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI-STREPTOCOCCAL/ANTI-MYOSIN IMMUNOGLOBULIN KAPPA
VARIABLE REGION (FRACMENT)).
Homo sapiens (Human).
Eukharyota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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Pred. No. 4.5e
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                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; Catarrhini; Hominidae;
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4.5e-05;
                                    DB 4;
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                                                         SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
Chen P., Deng J.B., Wang Z.L., Han H., Yao L.B.,
"Construction and sequencing of the single-chain
human TMF-alpha specific monoclonal antibody.";
human TMF-alpha specific monoclonal antibody.";
                                                                                                                                                                                                                                                                                                                      "Cloning genes of J. Cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               O9UL70
O9UL70;
O9UL70;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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Q9ERZ9;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI HUMAN TNF-ALDHA LIGHT CHAIN VARIABLE REGION (F)
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[3]
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Chen P.,
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Eukaryota; Metazoa; Chordata; Craniata; Ve
Mammalia; Eutheria; Primates; Catarrhini;
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Mammalia; Eutheria;
NCBI_TaxID=10090;
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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        SEQUENCE FROM N.A
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
Van der Merwe P.L.,
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l Similarity 94.4%;
17; Conservative
                                                                                                                                                                                                                                                                                                                          Deng J.B., Wang Z.L., Han H., Su C.Z. and sequencing of the light chain fra an anti-hTNF-a monoclonal antibody.", Mol. Immunol. 12:21-26(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                              FROM N.A.
Deng J.B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       108
108 AA;
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Rodentia;
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Pred. No. 4.6e
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Best Local Similarity 88.9
Matches 16; Conservative
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InterPro; IPR003006; Iq.
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InterPro; IPR003596; Iq.
Pfam; PF00047; Iq; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGy; 1.
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SEQUENCE 107 AA; 1178
Q9UL79;
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Q1-MAY-2000 (TrEN
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Q1-DEC-2001 (TrEN
Q1-DEC-2001 (TrEN
MYOSIN-REACTIVE )
(FRAGMENT);
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
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01-MAY-2000 (TIEMBLrel. 13, Created)
01-MAY-2000 (TIEMBLrel. 13, Last sequence update)
01-DEC-2001 (TIEMBLREL. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN KAPPA CHAIN VARIABLE
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NON_TER
SEQUENCE
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                         SMART;
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HSSP, P80362; IWTL.
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                                                                                                                                                                                                                                                                                                                                              "Myosin-reactive autoantibodies
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SM00406; IGV; 1.
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109
          (TrEMBLrel. 13, Created)
(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
TIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                  PRELIMINARY;
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11761 MW;
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                                                                                                                                                                                                                       FB1E43E7C7AFACCC CRC64;
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                                                                                                                                                                                                                                                                                                                                            in rheumatic
                                                                                                                                                                                                                                                                                                                                                                 Kalis N.N.,
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                                                                                                                                                                            9.4e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9.2e-05;
                                                                                                                                                                                        DB 4;
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                                                                                                                                                                                                                                                                                                                                            carditis
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                                                                                                                                                                                      Length 109
                                                                                                                                                                                                                                                                                                                                                                 Berney
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            REGION
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Best Local S
Matches 16
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NON_TER
SEQUENCE
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOSLOBULIN LIGHT CHAIN VARIABLE
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NON_TER
SEQUENCE
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EMBL; AF035031; AAI
HSSP; P80362; 1WTL
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
The Van der Merwe P.L., Kalis N.N.,
                                                                                                                                                                                                                                                    Interpro; IPA003006; Ig_MHC.
Interpro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
MCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clin. Immunol. Immunopathol. EMBL; AF035035; AAD56271.1; HSSP; P01607; 1REI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; 1g; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo Sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID-9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fetus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Myosin-reactive autoantibodies in rheumatic carditis and normal
                                                                                                                                                                                                                                                                                                                                                                                                                etus.";
                                                                                                                                                                                                                                                                                                                                                                                                                              Myosin-reactive autoantibodies in rheumatic carditis
                                        12
                                                                                                        Local Similarity
  65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12
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mes 16; Conserv
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SGSGTEFTLTISSLQFED
                    SGSGTDFTLTISSLQAED
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                                                                                      16;
                                                                                                                                                                                     108
108 AA;
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108 AA;
                                                                                   Conservative
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                                                                                                                                                                                         108
11834 MW;
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11787
                                                                                                        53.5%;
88.9%;
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88.9%;
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Pred. No.
                                                                                 Score 77; DB 4;
Pred. No. 0.00027;
l; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                       87:184-192(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                         9F9C5A92EBA96EEA CRC64;
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                                                                                                                            Length 108;
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                                                                                 Indels
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Best Local
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Q9UL78;
Q1-MAY-2000
                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.

MEDILINE-20250927; PubMed-10788485;

Radder C., Ritter G., Nathan S., Elia M., Gc
Cohen L.S., Welt S., Old L.J., Barbas C.F.

"The rabbit antibody repertoire as a novel
of therapeutic human antibodies.";
J. Biol. Chem. 275:13668-13676(2000).

EMBL: AF245502; AF68449.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        O9NOW5;
O1-OCT-2000 (TrEMBLrel. 15, Created)
O1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI-HUMAN A33 LIGHT CHAIN VARIABLE REGION (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NON_TER
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(FRAGMENT).
Homo sapiens (Human).
Homo sapiens (Human).
Homo sapiens (Human).
Homo sapiens (Human).
                                                                                           NON_TER
NON_TER
SEQUENCE
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q9N0W5
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SMART; SM00406; IGv; 1.
                                                                                                                                                                                                SMART;
                                                                                                                                                                                                                        InterPro; IPR003596; Pfam; PF00047; 19; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR003006; InterPro; IPR003596;
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                                                                                                                                                                                                                                                                                                                          HSSP; P80362; 1WTL
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                                                                                                                                                                                                SM00406;
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109 AA;
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                                                                                                                                                                                             IGV;
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11323
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                                                                                                                                                                                                                                                                                            Ig_MHC.
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                                                                                                  MW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 74; DB 4;
Pred. No. 0.0008;
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gout I.,
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Length 109;
                                                                                                  CRC64;
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Query Match

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Best Local
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01-OCT-2000
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                  NON_TER
NON_TER
SEQUENCE
                                                                                                              09JL82;
01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI-MYOSIN IMMUNOGLOBULIN LIGHT CHAIN VARIABLE REGION
     MEDLINE-20448942; PubMed-10992488; MEDLINE-20448942; PubMed-10992488; Malkiel S., Liao L., Cunningham M.W., Diamond B.; "T-Cell-dependent antibody response to the dominant epitope of streptococcal polysaccharide, N-acetyl-glucosamine, is cross-rewith cardiac myosin."; uith cardiac myosin."; Infect. Immun. 68:5803-5808(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIJINE-20448942; PubMed-10992488; Malkiel S., Liao L., Cunningham M.W., Diamond B.; "T-Cell-dependent antibody response to the dominant streptococcal polysaccharide, N-acetyl-glucosamine, with cardiac myosin."; Infect. Immun. 68:5803-5808(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI-MYOSIN IMMUNOGLOBULIN LIGHT CHAIN VARIABLE REGION
                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_V.
Pfam; PF00047; ig; 1.
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                                                                            STRAIN-BALB/C
                                                                                       SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=10090;
                                                                                                            NCBI_TaxID=10090;
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 AF206024; AAF69322.1;
                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                   99 AA; 10939 MW;
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Pred.
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Sciurognathi; Muridae;
                                                                                                                       Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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No.
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o the dominant
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                                cross-reactive
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Best Local S
Matches 14
Q99M37 PRELIMINARY; PRT; 238 AA.

Q99M37;
Q99M37;
Q1-JUN-2001 (TrEMBLrel. 17, Created)
T 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
T 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
T 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
E HYPOTHETICAL 26.3 KDA PROTEIN.

S Mus musculus (Mouse).
S Mus musculus (Mouse).
C Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
C Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
X NCBI_TaxID=10090;
N [1]
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Best Local S
Matches 14
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InterPro; IPR003006; Ig_N
InterPro; IPR003596; Ig_N
InterPro; IPR003596; Ig_N
Pfam; PF00047; 1g; 1.
SMART; SM00406; IGV; 1.
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Q9UL80;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Myosin-reactive autoantibodies fetus.";
Clin. Immunol. Immunopathol. 87:
EMBL; AF035034; AAD56270.1; -.
HSSP; P80362; IWTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (1)
SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed=9614934;
MEDLINE-98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON_TER
SEQUENCE
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR003006;
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rPro; IPR003596; Ig_v

n; PF00047; ig; 1.

RT; SM00406; IGv; 1.
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114 AA;
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104 AA;
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12775 MW;
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11360 MW;
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Primates;
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77.8%;
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2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 70;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87:184-192(1998)
                                         Craniata; Vertebrata; I
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Catarrhini; Hominidae;
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0.0032;
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                                         Euteleostomi; ; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Euteleostomi;
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RESULT
QPIWS9
ID WS9
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AC Q9
AC Q9
DT 011
DT 011
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DT 012
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Best Local S
Matches 13
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Best Local (
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InterPro; IPR003597; Ig_cl.
InterPro; IPR003600; Ig_like.
InterPro; IPR003600; Ig_MHC.
InterPro; IPR003596; Ig_V.
Pfam; PF00047; Ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 1.
SMARR; SM00400; IGv; 1.
SMARR; SM00410; IG_like; 1.
   time:
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Q91WS9;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
Q1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYPOTHETICAL 25.8 KDA PROTEIN (FRAGMENT).
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (SEP-2001) to the EMBL; BC013496; AAH13496.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Strausberg R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical protein.
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Hypothetical protein
SEQUENCE 238 AA; ;
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completed: July ne: 1484 sec
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14.104 Million cell updates/sec
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Copyright (c) 1993 - 2000 Compugen
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AAY34311	AAY34314	AAW10545	AAW10544	AAW10547	AAB03714	AAY50690	AAY05264	AAW21934	AAP80894	AAB07966	AAY18118	AAY18126	AAY23771	AAY23779	AAR70202	AAR75355	AAW06443	AAB78867	AAY77598	AAY57181	AAG65565	AAR61240	AAW27546	AAW15537	AAR45605	AAR30144	AAB62769	AAB18881	AAB18879	AAB18873	AAB18871	AAB18869	AAB18863
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ALIGNMENTS

AAW62807 RESULT

AAW62807 standard; Peptide;

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AAW62807;

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Green L,
Mendez M;
                                                                                                               03-DEC-1997;
                                                                                                                                                                                              Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa; production; antibody.
New transgenic non-human mammals - having an inactivated
                    WPI; 1998-333314/29
                                                                                           03-DEC-1996;
                                                                                                                                    11-JUN-1998.
                                                                                                                                                        WO9824893-A2
                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                             Amino acid sequence of a human antibody fragment.
                                                                                                                                                                                                                                                                                23-SEP-1998 (first entry)
                                                                      (ABGE-) ABGENIX INC.
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Best Local :
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New transgenic non-human mammals - having an inactivated aimmunoglobulin locus and a near complete human immunoglobulin locus, used for production of human antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
                                                                                                                                    WPI; 1998-333314/29
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                                                                                                                                                                                                                                                                                                    (ABGE-) ABGENIX INC.
                                                                                                                                                                                                                                                                                                                                                                 03-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                production; antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amino acid sequence of a human antibody fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                respectively.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 78; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunoglobulin locus and a near complete human immunoglobulin locus, used for production of human antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  52 sgsgtdftltissiqaed 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12 SGSGTDFTLTISSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 Similarity
18; Conserv
                                                                                                                                                                                                                                   Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ₹
                                                                                                                                                                                                                                                                                                                                                                    96US-0759620
                                                                                                                                                                                                                                                                                                                                                                                                                                     97WO-US23091.
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                                                                                                                                                                                                                                   Klapholz S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 87; DB 19;
Pred. No. 5.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                   Kucherlapati R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 82;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    locus, so that the mammal does not display normal B-cell development. The modified genome also has an inserted human heavy chain Ig locus in germline configuration, the human heavy chain Ig locus comprising a human micro constant region and regulatory and switch sequences, human J-H genes, human D-H genes, and human V-H genes and an inserted human kappa light chain Ig locus comprising a human kappa constant region, J-kappa genes, and V-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The transgenic animals have a near complete human Ight cous, including both a human heavy chain locus and a human kappa light chain locus. They can be used for the production of human antibodies when exposed to particular antigens e.g. when exposed to human II-8, EGFR or TNF- alpha tenestically.
                                                                                                                                                                                                                                                                                                                                                                                              Antibody 24-31; variable light chain; humanised antibody; gene therapy; CD40 binding inhibitor; gp39; autoimmune disease; food-related allergy; graft-versus-host disease; graft rejection; intestinal inflammation; reversible obstructive airways disease; allergy; transplant rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic Xenomice, created using the mammal which has genome specification describes a transgenic non-human mammal which has genome modifications that comprise an inactivated endogenous immunoglobulin (Ig) normal 8-cell development. The
Claim 7; Page 84; 122pp; English
                                New humanized antibodies to human gp30
                                                                                               Black A,
                                                                                                                                                                  08-SEP-1997;
                                                                                                                                                                                                  08-SEP-1998;
                                                                                                                                                                                                                                                                  WO9912566-A1
                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                               Mus sp.
                                                                                                                                                                                                                                                                                                                                                                immune response inhibitor; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antibody 24-31 humanised VL chain sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY05269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY05269 standard; Protein; 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW62793-822 represent fragments of human antibodies produced by transgenic Xenomice, created using the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 79; 128pp; English
                                                                                                                                 (IDEC-) IDEC PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                  asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          respectively.
                                                                                                                                                                                                                                 18-MAR-1999
                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                  leukaemia;
                                                                                                 Hanna N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A,
                                                                                                                                                                  97US-0925339
                                                                                                                                                                                                  98WO-US18163
                                                                                                 Newman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60.4%; Score 87; DB 100.0%; Pred. No. 6. Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                 immunosuppressant;
                                                                                                   Padlan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ž
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 19;
. 6.4e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 94;
                                                                                                                                                                                                                                                                                                                                                                                  cellular therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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RESULT
AAR52059
ID AAR52059
ID AAR52059
AC AAR5
XX AR5
AC AAR5
XX Anti
KW Iram
KW mode
OX Homc
XX Homc
XX Homc
XX Regi
FT Regi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence is the variable light chain of a humanised antibody of CC the invention. The antibody is capable of competing with a murine 24-31 CC antibody for inhibiting CD40 binding to gp39. The humanised antibodies CC can be used for the treatment of a disease treatable by modulating gp39 CC treating e.g. autoimmune diseases such as rheumatoid arthritis, psoriasis CC multiple sclerosis, diabetes, systemic lupus erythematosus and idiopathic CC multiple sclerosis, diabetes, systemic lupus erythematosus and idiopathic CC thrombocytopenic purpura (ITP) or non-autoimmune conditions such as CC graft-versus-host disease (GVHD) or graft rejection. They can also be CC used for the treatment of reversible obstructive airways disease, Crohn's CC disease and ulcerative colitis) and food-related allergies (e.g. cointestinal inflammations and allergies (e.g. coeliac disease, Crohn's CC disease and ulcerative colitis) and food-related allergies (e.g. migraine, rhinitis and escema), transplant rejection, asthma, leukaemia, or lymphoma. The antibodies can also be used as immunosuppressants, in CC particular during gene or cellular therapy. They may be used to inhibit humoral and cellular immune responses against viral vectors, e.g. certroviral vectors or adenoviral vectors. The use of such antibodies should enable such cells or vectors to be administered repeatedly, which continue advances and diseases such as cancers and continue and collition of chronic diseases such as cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
      09-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody; humanised; murine; human; heavy chain; light; variable;
framework region; complementarity determining region; reshaping;
                                                             07-SEP-1993;
                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                      Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR52059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR52059 standard; Protein; 111
                                                                                                                                                                               EP592106-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                              Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modelling; surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 SGSGTDFTLTISSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     65 sgsgtdftltisslgaed 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variable region of human KV4B antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
      92US-0942245
                                                             93EP-0307051.
                                                                                                                                                                                                                                                                                                                               /note-
63..95
                                                                                                                                                                                                                                                                                                                                                                                      /note=
56..62
                                                                                                                                                                                                                                         /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= complementarity_determining_region_1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       residue; modify.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               = framework_region_1
"FR 1"
                                                                                                                                                                                                                                         "CDR 3"
                                                                                                                                                                                                                                                                                                "FR 3"
                                                                                                                                                                                                                                                                                                                                                 "CDR 2"
                                                                                                                                                                                                                                                                                                                                                                                                                  "FR 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "CDR 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 87;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2v,
o. 7.4e-05;
o;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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AAW27527
ID AAW2
XX AAW2
XX 16-D
XX 16-D
XX Ligh
XX Gomp
KW Gomp
KW Inigh
KW Inigh
KW Chim
OS Home
XX JP09
XX JP09
XX 19-J
XX 19-J
XX 19-J
XX (ASP
PA (ASP
PA (TSU
XX Huma
PT high
PT anti
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                       (ASAK ) ASAHI
(NIKK-) NIKKA I
(TORI ) TORII
(TSUR/) TSURA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modification of a rodent antibody (Ab) or fragment by resurfacing in order to produce a humanised rodent Ab can be determined by calculating homology between murine and human Ab antibody surfaces. In order to test the resurfacing approach of the invention, three humanisation experiments were set up: (1) traditional loop grafting: (2) resurfacing approach using most similar chain; and (3) resurfacing approach using human sequences with most similar surface residues. The Ab used was the murine anti-N901 Ab (see AARS2055). Experiment 3 was carried out using the present sequence which represents the human KV4B Ab light chain variable region with 71 percent homology with anti-N901 Ab. N901/KV4B (AARS2060) was prepd. by resurfacing.
Humanised, semi-chimeric and chimeric antibodies against human high-affinity IgE receptor - useful medicinally and have low antigenicity in humans
                                                             WPI; 1997-429186/40.
                                                                                                                                                                                               19-JAN-1996;
                                                                                                                                                                                                                            29-JUL-1997
                                                                                                                                                                                                                                                         JP09191886-A
                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                    chimeric; treatment;
                                                                                                                                                                                                                                                                                                                              Complementarity determining region; CDR; murine; mouse; human; high affinity; immunoglobulin E; receptor; monoclonal antibody; light chain; variable region; humanised; semi-chimeri
                                                                                                                                                                                                                                                                                                                                                                                            Light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW27527 standard; Protein; 112 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method of resurfacing of rodent antibodies to produce humanised antibody forms - for producing non-human antibodies with improved therapeutic efficiency by presenting human surface on V-region
                                                                                                                                                                                                                                                                                                                                                                                                                          16-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW27527;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Guild BC,
                                                                                                                                                                  19-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Fig 4A; 230pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1994-120230/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PEDE/) PEDERSEN J T. (IMMU-) IMMUNOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pedersen JT,
                                                                                                                                                                                                                                                                                                                                                                                            variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                         YAKUHIN
T.
                                                                                                                       WHISKEY
                                                                                                                                     BREWERIES LTD
                                                                                                                                                                 96JP-0024816
                                                                                                                                                                                               96JP-0024816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60.4%;
                                                                                                                                                                                                                                                                                                                 prevention; disease; allergy; CRA2
                                                                                                                                                                                                                                                                                                                                                                                         region of human CRA2 antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rees
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 87;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 15;
7.7e-05;
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                                                                                                                                                                                                                                                                                                                               semi-chimeric;
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RESULT
ADARSO313
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AARSO313
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AARSO313
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AARSO313
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AARSO32
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AARSO32
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monoclonal antibody; Plasmodium falciparum; CDR; complementarity determining region; fusion protein; murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                                                        (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                            Claim 5;
                                                                                                                                                                                                                                      Charoenvit Y,
Sadoff JC, Sy
                                                                                                                                                                                                                                                                                                                                                                                        09-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09405690-A
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Naturally-occurring (AAQ44841-42)
                                                                                                        New engineered antibodies and fusion Plasmodium infection - contg. murine
                                                                                                                                                                          N-PSDB;
                                                                                                                                                                                              WPI; 1994-101115/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                08-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         69 sgsgtdftltisslqaed 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2;
                                                                                                                                                                                                                                                                                                 SMITHKLINE BEECHAM CORP
US SEC OF ARMY.
US SEC OF NAVY.
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                                          Fig
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                                                                                      nucleic acid, vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     112
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                                                                                                                                                                                                                                      Sylvester
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                          2;
                                                                                                                                                                                                                                      Hoffman S,
lvester DR,
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                                       98pp;
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95..10
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24..40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label
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                                            English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - CDR2
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Pred. No.
                                                                                        and transformed
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  and synthetic (AAQ44825-28)
                                                                                                          proteins
antibody
                                                                                                                                                                                                                                                              Rosenberg
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7.8e-05;
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                                                                                                        r preventing
R sequences,
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb NFS2, its variable chain peptides, CDRs, functional fragments, rand analogs are useful in prodn. of fusion proteins, esp. engineered antibodies. These prods. are used to protect humans against Plasmodium infections.
                                                                                                                                                                                                     (SMIK )
(USSA )
(USNA )
                                                                                                                                                                                                                                                                                                                                                                                                                                           complementarity determining region; fusion protein; murine; variable; light; heavy; chain; malaria.
Naturally-occurring (AAQ44841-42) and synthetic (AAQ44825-28) variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb NFS2, its variable chain peptides, CDRs, functional fragments, Fab fragments, and analogs are useful in prodn. of fusion proteins, esp. engineered antibodies. These prods. are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR50314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                         Claim
                                                                                                 New engineered antibodies and fusion proteins for preventing plasmodium infection - contg. murine antibody CDR sequences, corresp. nucleic acid, vectors and transformed cells
                                                                                                                                                                        Charoenvit Y, Hoffman Sadoff JC, Sylvester
                                                                                                                                                                                                                                                  09-SEP-1992;
                                                                                                                                                                                                                                                                     08-SEP-1993;
                                                                                                                                                                                                                                                                                          17-MAR-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Humanised light chain variable
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18; Conser
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                                                                                                nucleic acid,
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56..62
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95..103
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100.0%; Pr
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DR,
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Gross
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                                                                                                                                                                         MS;
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                                                                                                                                                                                     Rosenberg
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3. 7.8e-05;
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Best Local
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                                                              Ames RS, Ar
                                                                                                                   06-JUN-1995;
23-DEC-1994;
06-JUN-1995;
                                                                                          (SMIK )
     New monoclonal antibody to human interleukin-5 - used to produce products for the treatment and diagnosis of conditions associated with excess eosinophil prodn., e.g asthma etc.
                                                                                                                                                                                                                                                                                                                                                                           interleukin-5; IL-5; eosinophil; asumma, uniterleukin-5; IL-5; eosinophil; asumma, uniterleukin-5; therapy; diagnosis; light chain; VL; monoclonal antibody; MAb.
                                         N-PSDB; AAT34094
                                                                                                                                                                      11-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Humanised
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                                                  1996-333976/33.
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18; Conservative
                                                                                           SMITHKLINE SMITHKLINE
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                                                                 Appelbaum ER, McMillan LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                286
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                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                              antibody light chain variable region.
                                                                                                                   95US-0470110.
94US-0363131.
95US-0467420.
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63..94
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56..62
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41..55
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                                                                                                                                                                                                       /note-
                                                                                                                                                                                                                                                                                                                                  /note= "human LEN framework region"
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                                                                                           BEECHAM CORP
BEECHAM PLC.
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100.0%;
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                                                                Chaiken IM,
Theisen TW;
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Pred. No.
                                                                                                                                                                                                                                                LEN
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                                                                                                                                                                                                                                                                                                          286
                                                                                                                                                                                                      framework region"
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                                                                          Cook RM,
                                                                                                                                                                                                                                                                                                          VL CDR1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 15,
J. 7.8e-05;
O;
                                                                         Gross MS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 113;
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RESULT
AAW42466
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Best Local S
Matches 18
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                                                    24 - DEC - 1997
                                                                         WO9748418-A1
                                                                                                                   Region
                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                                 Chimeric -
                                                                                                                                                                                                                                                                                                                                                                              Neutralising antibody; monoclonal antibody; MAb; 2B6; LEN; mouse; interleukin-5; IL-5; human; eosinophil; asthma; allergic rhinitis; atopic dermatitis; therapy; diagnosis; humanised antibody.
                              20-JUN-1997;
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                                                                                                                                                                                                                                                                                                                 Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 55-56; 120pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              113 AA;
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                                                                                                                                                                                                                                                                                                                                                           musculus
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100.0%; Pr
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Pred. No.
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7.8e-05;
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Best Local
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   Polynucleotides encoding anti-p53 antibodies, polypeptides and fragments, useful in treatment and diagnosis of cancer, rheumat
                                                                                       N-PSDB; AAA96134.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200056770-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino acid sequence of anti-p53 antibody light chain clone 163.2.
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                                                                                                                  WPI; 2000-638249/61.
                                                                                                                                                                                                                                 (SVIN-)
                                                                                                                                                                                                                                                                                             19-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                    15-MAR-2000; 2000WO-AU00189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   p53; antibody; immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71 sgsgtdftltisslqaed 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 SGSGTDFTLTISSLQAED 29
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                                                                                                                                                                                                                                    ST VINCENT'S HOSPITAL SYDNEY LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      113 AA;
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                                                                                                                                                                                                                                                                                          99AU-0009321
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   response; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English
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Pred. No.
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7.8e-05;
ches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies care useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a cc vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as cc well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis can properticular, the following can be treated cancer, rheumatoid arthritis cf and coronary heart disease. Cancer include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, chead and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, breast cancer, brain cancer, bladder cancer, prostate cancer and curinary/genital tract cancer, oesophageal cancer, mesenchymal tumours, cc e.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma.
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Best Local
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                                                                                                                                                N-PSDB;
                                                                                                                                                              WPI; 2000-638249/61.
                                                                                                                                                                                            Ward RL,
                                                                                                                                                                                                                                                                                                               28-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                   p53; antibody; immune response; vaccine; gene therapy;
rheumatoid arthritis; coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amino acid sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-FEB-2001
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                                                                                                                                                                                                                        (SVIN-)
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                                                                                                                                                AAA96138
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                                                                                                                                                                                            Coomber DWJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and coronary heart disease
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100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 anti-p53 antibody light chain clone 163.6.
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                                                                                                                                                                                                                          SYDNEY LTD
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The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies

Claim

30; Page 136; 163pp; English.

arthritis and coronary heart disease

Polynucleotides encoding anti-p53 antibodies, fragments, useful in treatment and diagnosis arthritis and coronary heart disease

of cancer, rheumatoid

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RESULT 1
AAB1863
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XX P53;
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Best Local S
Matches 18
                           The present sequence represents the light chain of an antibody reactive against p33. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ward
                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                     Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease
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                                                                                                                                                                                                                                                                                           and coronary heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coomber DWJ;
                                                                                                                                                                                                                           Page 138; 163pp;
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The polypeptide components of the antibodies are useful
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Pred. No.
                                                                                                                                                                                                                                                                                              disease
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7.8e-05;
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           an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ward
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                                                                                                                                                                                                                                                                                       The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing
                                                                                                                                                                                                                                                                                                                                                                                                                            arthritis and coronary heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB18869 standard; Protein; 113 AA
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                                                                                                                                                                                                                                                                                                                                                                         Claim 30; Page 144; 163pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     and diagnosis
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7.8e-05;
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In particular, the following and coronary heart disease. of epithelial origin, e.g. (

colo-rectal

cancer,

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Best Local Similarity
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100.0%;
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Pred. No.
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7.8e-05;
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12

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Query Match
Best Local Similarity
Matches 18; Conserv

Conservative

0

60.4%;

Score 87; DB s; Pred. No. 7.8 0; Mismatches

DB 21; 7.8e-05

21;

Length 113

Indels

0;

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RESULT 1
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                           The present sequence represents the light chain of an antibody reactive cagainst p53. The antibody is obtained from a vertebrate host expressing can immune response against a naturally occurring disease. The antibodies considered the compositions of the antibodies considered the compositions of the antibodies considered the compositions of the antibodies are useful in pharmaceutical compositions, which additionally contain considered the considered the considered the considered the antibodies are useful in vaccines, for inducing an immune response against a disease in a convertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as convertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as convertebrate, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancer include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, chead and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, castric cancer, brain cancer, cancer and cancer, brain cancer, brain cancer, and hemopoletic tumours, e.g. B cell lymphoma.
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Best Local S
Matches 18
Sequence
                                                                                                                                                                                                                                                                                                                                                                              Claim 30; Page 148; 163pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                               fragments, useful in treatment and diarthritis and coronary heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                               Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               p53; antibody; immune response; vaccine; gene therapy; cancer;
rheumatoid arthritis; coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino acid sequence of anti-p53 antibody light chain clone 163.17
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100.0%; Pr
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7.8e-05;
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71 sgsgtdftltisslqaed 88

Search completed: July 15, 2002, 12:57:59 Job time: 415 sec

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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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Match
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/2/iaa/Dackfiles1.pep:*
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144
1 YERKKRRQRRRSGSGTDFTLTISSLQAED
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US-08-554-840-13
US-08-554-840-15
US-08-554-840-25
US-08-525-39A-81
US-07-942-245-30
US-08-467-769A-21
US-08-467-769A-21
US-08-637-769A-21
US-08-637-647-21
US-08-525-539A-80
PCT-US93-08435-6
PCT-US93-08435-6
PCT-US93-08435-6
US-08-450-578-6
US-08-050-5769B-31
US-08-050-769B-31
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sequence
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Sequence
Sequence
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13, Appl
2, Appl
30, Appl
21, Appl
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16, Appl
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11, Appl
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SGSGTDFTLTISSLQAED 29

Matches

18;

Conservative

0;

Mismatches

0

Indels

0;

Gaps

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Query Match 60.4%; Score 87; DB 3; Length 80; Best Local Similarity 100.0%; Pred. No. 1.1e-05;	US-08-54-840-13 US-08-554-840-13 US-08-554-840-13 Sequence 13, Application US/08554840 PAtent No. 6001358 GENERAL INFORMATION: APPLICANT: HANNA, Nabil APPLICANT: HANNA, Nabil APPLICANT: PADLAN, Eduardo A. APPLICANT: NEWNAN, Roland A. APPLICANT: NEWNAN, Roland A. ITILE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPEUT! NUMBER OF SEQUENCES: 28 CORRESPONDENCE ADDRESS: ADDRESSEE: Burns, Doane, Swecker & Mathis STREET: PO. Box 1404 CITY: Alexandria STATE: VIGANIa STATE: VIGANIa COUNTRY: United States COUNTRY: United States COUNTRY: Onited States APPLICATION TYPE: FLOPPY disk COMPUTER IBADABLE FORM: MEDIUM TYPE: PLOPPY disk COMPUTER: IBADABLE FORM: MEDIUM TYPE: PLOPPY disk COMPUTER: SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION NUMBER: US/08/554,840 FILING DATE: 07-NOV-1995 CLASSIFICATION NUMBER: 35,030 REFERENCE/DOCKET NUMBER: 35,030 REFERENCE/DOCKET NUMBER: 012712-127 TELECOMMUNICATION HORDMATION: TELEFAX: (703) 836-6620 TELEFAN: (703) 836-6620 TELEFONE: CHARACTERISTICS: LENGTH: 80 amino acids TYPE: amino acid STRANDEDWESS: ilnear SOFTSH HOLDWESS: ilnear MOLECULE TYPE: protein	ALIGNMENTS	28 87 60.4 133 3 US-08-463-903-4 Seq 29 87 60.4 133 4 US-07-935-695-4 Seq 30 87 60.4 135 3 US-08-812-518-14 Seq 31 87 60.4 155 3 US-08-812-718-11 Seq 32 87 60.4 171 3 US-09-160-567-11 Seq 33 87 60.4 171 4 US-07-935-695-20 Seq 34 87 60.4 241 2 US-07-916-098A-56 Seq 36 87 60.4 260 3 US-08-463-903-2 Seq 37 87 60.4 260 3 US-08-463-903-2 Seq 38 87 60.4 275 3 US-08-463-903-2 Seq 40 87 60.4 275 3 US-08-463-903-2 Seq 41 87 60.4 285 3 US-08-463-903-2 Seq 42 87 60.4 285 4 US-07-935-695-2 Seq 42 87 60.4 285 4 US-07-935-695-2 Seq 43 87 60.4 342 3 US-08-828-741B-6 Seq 44 87 60.4 342 3 US-08-828-741B-6 Seq 45 87 60.4 495 3 US-08-628-741B-4 Seq
	THERAPEUTIC USE THEREOF		Sequence 4, Appli Sequence 46, Appli Sequence 11, Appli Sequence 11, Appli Sequence 20, Appli Sequence 20, Appli Sequence 20, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 22, Appli Sequence 22, Appli Sequence 22, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 4, Appli

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                                                                                                                               GENERAL INFORMATION:
APPLICANT: BLACK, I
APPLICANT: HANNA, I
APPLICANT: PADLAN,
                                                                                                                                                                                                     Sequence 2, Application US/08554840 Patent No. 6001358
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Best Local :
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FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 0127
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                    APPLICANT: NEWMAN, ROLAND A.
TITLE OF INVENTION: HUMANIZED ANT
TITLE OF INVENTION: COMPOSITIONS
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Burns, Doane, Swecker & Mathis
STREET: P.O. Box 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: NEWMAN, ROLAND A.
TITLE OF INVENTION: HUMANIZED ANT
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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APPLICANT: HANNA, Nabil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
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les 18; Conserv
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                                                          NEWMAN, COLAND A.

NEWMAN, ROLAND A.

VENTION: HUMANIZED ANTIBODIES TO HUMAN 9p39,

COMPOSITIONS CONTAINING AND THERAPE
                                                                                                                                                HANNA, Nabil
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                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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100.08; Pr
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HUMANIZED ANTIBODIES TO HUMAN 9D39,
COMPOSITIONS CONTAINING AND THERAPI
                                                                                                                             Eduardo A.
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                                                                           AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                       Length 80;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-525-539A-81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 81, Application US/08525539A Patent No. 6309636
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
FILING DATE: 14-SEP-1995
CLASSIFICATION: 424
ATTORNEY_AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,612
REFERENCE_POCKET NUMBER: 2763
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION UMBER: US/08/554
FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                              STREET: /JC
CITY: Palo Alto
CTMATE: CA
                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESSE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: DO COUTO, FERNANDO J.R. APPLICANT: CERIANI, ROBERTO L. APPLICANT: PETERSON, JERRY A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 94304-1018
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65 SGSGTDFTLTISSLQAED 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                  RECOMBINANT PEPTIDES DERIVED FROM THE MC3 ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF, METHODS OF HUMANIZING ANTIBODY PEPTIDES
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100.0%;
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                                                                                                                       US/08/525,539A
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                     27633-20001.21
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Pred. No.
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                                                                                                                                                        Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DВ 3;
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;; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-07-942-245-30
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RESULT 6
US-08-467-420A-21
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                                                                                                                                                                                                                                                                              TELECOMMUNICATION: 530
TELECOMMUNICATION: TORRATION: TELEPHONE: (202) 293-7060
TELEPHONE: (202) 293-7860
TELEPAX: (202) 293-7860
TELEPAX: (202) 293-7860
TELEPAX: 6491103
INFORMATION FOR SEQ ID NO: 30:
                                                                                                                Query Match
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local S
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TELEX: 706141
INFORMATION FOR SEQ ID NO: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ROGUŚKA, MI
APPLICANT: GUILD, Br;
TITLE OF INVENTION: SUR!
TITLE OF INVENTION: ANT
TITLE OF SEQUENCES: 52:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTER:
2019: 20037-3202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy Workstation
COMPUTER: HP 9000/700 Workstation
                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: In house
CURRENT APPLICATION DATA:
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TYPE: amino acid
STRANDEDNESS: single
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LENGTH: 107 amino acids
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                                                                          12 SGSGTDETLTISSLQAED 29
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OPERATING SYSTEM:
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CITY: Washington
STATE: D.C.
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                                                         SGSGTDFTLTISSLQAED 88
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5639641
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18; Conservative
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2100 Pensylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEARLE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anthony R.
Anthony R.
A, Michael A.
Braydon C.
SURFACE RESIDUE VENEERING OF ANTIBODIES
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100.0%; Pred. No.
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                                                                                                                  100.0%; I
tive 0;
                                                                                                                                60.4%; Score 87; DB 1; I 100.0%; Pred. No. 1.5e-05;
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                                                                                                                   Mismatches
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                                                                                                                                            Length 112;
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US-08-470-110A-21

; Sequence 21, Application US/08470110A

; Patent No. 5693323
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; MOLECULE TYPE:
US-08-467-420A-21
Sequence 21, Application of the control of the cont
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Best Local S
Matches 18
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
TOMPUTER: IBM PC compatible
TOMPUTER: PC-DOS/MS-DOS
TOMPUTER: TOMPUTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 610 270-5090 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EILING DATE:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34 02
REFERENCE/DOCKET NUMBER: P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Smithkline Beecham Corp./Corporate
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APPLICANT:
APPLICANT:
APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinan
TITLE OF INVENTION: Treatment
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APPLICANT: Appelbaum, Edward
APPLICANT: Chaiken, Irwin M.
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CITY: King of Prussia
STATE: Pennsylvania
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18; Conserv
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Cook, Richard M. Gross, Mitchell S. Holmes, Stephen D.
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Holmes, Stephen D.
                                                                                                        Appelbaum, Edward Chaiken, Irwin M.
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                                                                                                                                                                         Ames, Robert S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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100.0%; Pred. No. 1.6e-05;
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APPLICANT:

McMillan, Lynette J Theisen, Timothy W.

NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS: TITLE OF INVENTION:

Recombinant IL5 Antagonists Useful in Treatment of IL5 Mediated Disorders

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                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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Best Local :
                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 610 270-5090 INFORMATION FOR SEQ ID NO: 21:
                                                                                                           APPLICANT: McMillan, Lynette J.
APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
TITLE OF INVENTION: 76
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LENGTH: 113 amino acids
TYPE: amino acid
STRANDEDNESS: single
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TELECOMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
TELEPAX: 610 270-5024
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APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PATENTIN ReLease #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                      NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEB: SmithKline Beecham Corp./Corporate
STREET: P.O. Box 1539-UW2220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
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COMPUTER READABLE FORM:
                                                      CITY: King of Prussia
STATE: Pennsylvania
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les 18; Conservative
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                  COUNTRY: USA
ZIP: 19406-0939
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            71 SGSGTDFTLTISSLQAED 88
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                                                                                                                                                                                                                                                                                                                                                                  21, Application US/08667769A
o. 5783184
                                                    Pennsylvania
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                                                                                                                                                                                                                                    Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
                                                                                                                                                                                                                                                                                           Appelbaum, Edward R. Chaiken, Irwin M.
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100.0%; pr
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pred. No. 1.6
0; Mismatches
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hes 0;
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                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Ames, R
APPLICANT: Appelba
APPLICANT: Chaiken
                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21, Application US/08940371 patent No. 5851525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local Similarity
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/467420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                           NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp./Corporate
ADDRESSEE: Intellectual Property
                                                                                                                                                                                                                                                                                         APPLICANT:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER: PCT/
FILING DATE: 22-DEC-1995
                                                                                                                                                                                                 APPLICANT: McMillan, Lynette J.
APPLICANT: Thelsen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                                                                          APPLICANT:
                                     COUNTRY:
ZIP: 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 SGSGTDFTLTISSLQAED 29
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                                                                                       STREET: P. O. Box 1539-UW2220
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            71 SGSGTDFTLTISSLQAED 88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/667,769A
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                                         19406-0939
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                                                                        Pennsylvania
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                                                                                                                                                                                                                                                                          Gross, Mitchell S. Holmes, Stephen D.
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                                                                                                                                                                                                                                                                                                             Cook, Richard M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21,
Patent No. 6
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Best Local
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PILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
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                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in TITLE OF INVENTION: Treatment of IL5 Mediated Disorders NUMBER OF SEQUENCES: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
                                                 PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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CITY: King of Prussia
STATE: Pennsv'"...'
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                                                                 APPLICATION NUMBER: US/08/637,647 FILING DATE: 19-JUN-1997 CLASSIFICATION: 536
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STRANDEDNESS: sin
TOPOLOGY: linear
CLASSIFICATION:
                                 APPLICATION NUMBER:
                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 19406-0939
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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                               US/08/363,131
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; Pred. No. 1.6e-05;
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US-08-525-539A-80
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Best Local Similarity
Matches 18; Conserv
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                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 610 270-5090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                              APPLICATION NUMBER: US/01
FILING DATE: 14-SEP-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
                                                                                                                                            REGISTRATION NUMBER: 37,612
REFERENCE/DOCKET NUMBER: 7,71
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: DO COUTO,
APPLICANT: CERIANI, F
APPLICANT: PETERSON,
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REFERENCE/DOCKET NUMBER: 95
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
                                                                               SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
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NAME: Sutton, Jeffrey A.
                                                                                                                TELEPHONE: (415) --- TELEPHONE: (415) 494-0792
              STRANDEDNESS:
TOPOLOGY: li
                                               TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: MUKKLOU. STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94304-1018
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              linear
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                             single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JERRY A.
                                                                                                                                                                                                                                                                                    US/08/525,539A
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Query Match Best Local Similarity

60.4%; 100.0%;

Score 87; Pred. No.

DB 4; I 1.6e-05;

Length 113;

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PCT-US93-08435-6
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PCT-US93-08435-8
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (215) 540-5818 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acids
TOPOLOGY: 115-
WOLECUTO 115-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/941,654
FILING DATE: 09-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08435
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REFERENCE/DOCKET NUMBER: SBUTELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: NO
TITLE OF INVENTION: IM
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS: LENGTH: 113 amino acid
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                                   APPLICANT:
                                                   AFPLICANT:
                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                   APPLICANT:
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CITY: Spring House
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                                                                                                                                                                                                          71 SGSGTDFTLTISSLQAED 88
                                                                                                                                                                                                                                        12 SGSGTDFTLTISSLQAED 29
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OF INVENTION: Novel Antibodies for Conferring Passive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SmithKline Beecham, Corporation U. S. Government, Secretary of the Navy
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                              U. S. Government, Secretary of the Navy
U. S. Government, Secretary of
                                                                  SmithKline Beecham, Corporation U. S. Government, Secretary of
                 the Army
                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (215) 540-9200
                                                                                                                                                                                                                                                                                            60.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel Antibodies for Conferring Passive Immunity Against Infection by a Pathogen in Man
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PCT-US95-17082A-21
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                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21, Application PC/TUS9517082A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local :
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                          APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward
APPLICANT: Chalken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell (APPLICANT: Holmes, Stephen I APPLICANT: MCMillan, Lynette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION:
CRASSIFICATION DATA:
APPLICATION NUMBER: US 07/941,654
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MEDIUM TYPE: Floppy
   ZIP: 19406-0939

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                           NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                          APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 SGSGTDFTLTISSLQAED 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Howson and Howson STREET: Box 457, 321 Norristown Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71
                                                                                                                                       ADDRESSEE: SmithKline Beecham Corp./Corporate STREET: P.O. Box 1539-UW2220 CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100
les 18; Conservative
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                                                                                                         COUNTRY:
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Chaiken, Irwin M.
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100.0%;
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       Version #1.30
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TITLE OF INVENTION: Immunity Against Infection by a Pathogen in NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                     Version
                           DB 5;
               1.6e-05;
                           Length 113;
Indels
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 Gaps
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CÜRRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/17082A

PRIOR APPLICATION

CLASSIFICATION: FILING DATE:

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; TOPOLOGY: linear
; MOLECULE TYPE: protein
PCT-US95-17082A-21
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US-08-360-125-6
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: SULTON, Jeffrey A.
REGISTRATION NUMBER: 34 02
REFERENCE/DOCKET NUMBER: P
                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/905,
FILING DATE: June 29, 1992
                                                                                              SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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                                                      FILING DATE:
CLASSIFICATION:
                                                                                                                           COMPUTER: IBM Compatible OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                  CITY: Washington STATE: D.C.
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Local Similarity 100.08;
                                                                                  APPLICATION NUMBER:
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APPLICATION NUMBER:
                                                                                                                                                                                                  COUNTRY:
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TYPE: amino acid
STRANDEDNESS: single
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805 Fifteenth
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Kazuhiro NAGAIKE
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                                                                                                                                                                                                                                                                                                                                                                       Yoko HIRAKAWA
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RER: US_08/470110
                                                                                                                                                                                                                                                                                                   Specifically Binding to Surface Antigen of Cancer Cell Membrane
                                                                                                                                                                                                                                                                                                                              Human Monoclonal Antibody
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                         07/905,534
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Street, N.W., #700
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Pred. No.
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Search completed: July 15, Job time: 391 sec

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; FILING DATE:
; PUBLICATION DATE:
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US-08-360-125-6
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Best Local (
                                                    Matches
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                                                                                                                                                                                                                                               PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                          POSITION IN GENOME:
CHROMOSOME/SEGMENT:
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IMMEDIATE SOURCE:
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ORIGINAL SOURCE:
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REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
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                                                                                                                                                                                                                                                                                                                                                                                                                                  CELL TYPE:
CELL TYPE:
CELL LINE:
               12 SGSGTDFTLTISSLQAED 29
                                                              Local Similarity
                                                                                                                                                                                                            JOURNAL:
VOLUME:
                                                                                                                                                                                                                                                                                               NAME/KEY:
LOCATION:
71 SGSGTDFTLTISSLQAED
                                                                                                                                                                                                                                                                       IDENTIFICATION METHOD: OTHER INFORMATION:
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TISSUE TYPE:
                                                                                                                                                                          DATE:
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DEVELOPMENTAL STAGE:
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                                                Conservative
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linear
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100.0%;
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                                               Score 87; DB 1; ; Pred. No. 1.6e-05 0; Mismatches 0
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Title:
Perfect score:
Sequence:
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Maximum
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12 SGSGTDATAPASSLQAED

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Query Matc Best Local Matches	Ig kappa chain V region (V-kapp C;Species: Homo sapiens (man) C;Date: 06-Jan-1995 #sequence_r C;Accession: S37530 R;KLein, U: Kueppers, R.; Raje submitted to the EMBL Data Libr A;Description: Human IgM(+)IgD A;Reference number: S37501 A;Accession: S37530 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-92 <kle>A;Cross-references: EMBL: 226630 C;Superfamily: immunoglobulin VC;Keywords: heterotetramer; imm</kle>	Query Matches Amatches Qy 112 SG 111 Db 49 SG RESULT 2	A;Accession: S37533 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-92 <kle> A;Cross-references: EMBL:Z2 C;Superfamily: immunoglobul C;Keywords: heterotetramer;</kle>	RESULT 1 S75353 Ig kappa chain V regio C;Species: Homo sapien C;Date: 06-Jan-1995 #s C;Accession: S37533 R;Klein, U.; Kueppers, submitted to the EMBL A;Description: Human I A;Reference number: S3		4444433 330 4444 4421 543
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Score 63; DB 2; Pred. No. 0.31; ; Mismatches	- human (fragment) ion 06-Jan-1995 #text_ch K. September 1993 ells, the major B cell s ells, the major B cell s on; immunoglobulin homo	red. No. 0.31; Mismatches	05708; PI immunogl) - human (fragment sion 06-Jan-1995 #t y, K. , September 1993 cells, the major B	ALIGNMENTS	050 050 051 052 0054 0054 0053 0053 0053 0053 0053 0053
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Ig kappa chain V region (V-kappa 4) - human (fragment)
C:Species: Homo sapiens (man)
C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C:Accession: S37535
R:Klein, U.; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the periphe:
A;Reference number: S37501
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 <KLE>
A;Cross-references: EMBL:Z26625; NID:g405712; PIDN:CAA81378.1; PID:g405713
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
Ig kappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change:
C;Accession: S37531
R;Klein, U: Kueppers, R: Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset A;Reference number: S37501
A;Raccession: S37531
A;Status: preliminary
A;Molecule type: mRNA
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submitted to the EMBL Data Library, September 1993
A.Description: Human IgM(+)IgD(+) cells, the major B cell subset in the pe.
A;Reference number: S37501
A;Accession: S37529
A;Status: preliminary
A;Molecule type: mRNA
A.Residues: 1-92 <KLE>
A;Residues: 1-92 <KLE>
A;Cross-references: EMBL:Z26631; NID:g405700; PIDN:CAA81384.1; PID:g405701
C;Keywords: heterotetramer; immunoglobulin
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S37529
Ig kappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: S37529
R;Klein, U.; Kueppers, R.; Rajewsky, K.
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C;Accession: S37534
C;Accession: S37534
R;Klein, U.; Kueppers, R.; Rajewsky, K.
R;Klein, U.; Kueppers, Lata Library, September 1993
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peripheral blance number: S37501
                                                        Query Match
Best Local Similarity
Watches 14; Conserve
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R;Klein, U.; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset
A;Reference number: S37501
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A; Residues: 1-92 <KLES
A; Cross-references: EMBL: 226626; NID: 9405710; PIDN: CAA81379.1; PID: 9405711
C: Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
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$37534

Kappa chain V region (V-kappa 4) - human (fragment)

Ig kappa chain V region (man)

C;Species: Homo sapiens (man)

C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                       Ig kappa chain V region (V-kappa 4) - human (fragment)
C; Species: Homo sapiens (man)
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A; Residues: 1-92 <KLE>
                                                                                                                                                                                                                                                                            A; Status: preliminary
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Best Local
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                                     12 SGSGTDATAPASSLQAED 29
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SGSGTDFTLTISSLQAED
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77.8%;
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Pred. No. 0.31;
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Ig kappa chain V region (anti-DNA, H2F) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 21-Jan-2000
C;Accession: PH0869
C;Accession: PH0869
R;Manheimer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, B.
J. Exp. Med. 174, 1639-1652, 1991
A;Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idiotype
A;Reference number: PH0862; MUID:92078875
A;Accession: PH0869
A;Molecule type: DNA
A;Residues: 1-101 <MAN>
A;Residues: 1-101 <MAN>
A;Note: residues 28-33 were obtained from Figure 4
C;Comment: This antibody is produced by Epstein-Barr virus-transformed B cell that bears
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Reywords: heterotetramer; immunoglobulin
F;14-3/Region: framework 1
F;16-5/Deadin: complementarity-determining 1
F;41-55/Region: complementarity-determining 2
F;56-7/Region: complementarity-determining 2
A49138

IgA kappa rheumatoid factor variable - human C; Species: Homo sapiens (man)
C;Date: 19-Dec-1993 #sequence_revision 18-No C;Accession: A49138
R;Gause, A.; Kuppers, R.; Mierau, R.
Clin. Exp. Immunol. 88, 430-434, 1992
A;Title: A somatically mutated V kappa IV ge A;Reference number: A49138; MUID:92298590
A;Accession: A49138
A;Status: preliminary
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F;63-94/Region: 1
F;95-101/Region:
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A; Residues: 1-102 <MIH>
A; Residues: 1-102 <MIH>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology <IMM>
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| 71 SGSGTDFTLTISSLQAED 88
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Pred. No. 0.33;
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Ig kappa chain V region - hum:
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence
C;Accession: S30520
R;Mariette, X.
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A:Residues: 1-106 <GAU>
A:Residues: 1-106 <GAU>
A:Residues: 1-106 <GAU>
A:Residues: 1-106 <GAU>
A:Residues: GB:S37926; NID:g298207; PIDN:AAB22366.1; PID:g298208
A:Note: sequence inconsistent with the nucleotide translation
A:Note: sequence extracted from NCBI backbone (NCBIN:106633, NCBIP:106637
C:Superfamily: immunoglobulin V region; immunoglobulin homology
F;20-99/Domain: immunoglobulin homology <IMM>
                                                                                                                                 A;Cross references: EMBL:218325
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-113 <MAR>
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A; Status: preliminar
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C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;16-96/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-113 < MAR>
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A; Accession: S30523
A; Status: preliminary
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A; Reference number: S30520
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                                                            Similarity
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Pred. No. 0.35;
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                                        Score 63; DB 2;
Pred. No. 0.37;
0; Mismatches
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Pred. No. 0.37;
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Ig kappa chain V-IV region (Len) - human
C;Species: Homo sapiens (man)
C;Date: 24-Apr-1984 #sequence_revision 24-Apr-1984 #text_change 01-Dec-2000
C;Date: 24-Apr-1980 #sequence_revision 24-Apr-1984 #text_change 01-Dec-2000
C;Accession: A01903; F61458
R;Schneider, M.; Hilschmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 356, 507-557, 1975
A;Title: Die Primaerstruktur einer monoklonalen Immunglobulin-L-Kette der St.
A;Title: Die Primaerstruktur einer monoklonalen Immunglobulin-L-Kette der St.
A;Reference number: A01903; MUID:76004342
A;Accession: A01903
A;Accession: A01903
A;Accession: A01903
A;Molecule type: protein
A;Note: this is the first completely sequenced V region of a new kappa chain
A;Note: the C region of this chain has the Inv (3) marker
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C.Species: Homo sapiens (man)
C.Species: Homo sapiens (man)
C.Accession: S34003
R.Mariette, X.; Tsapis, A.; Brouet, J.C.
Eur. J. Immunol. 23, 846-851, 1993
A.;Title: Nucleotidic sequence analysis of the variable domains of four humar A; Reference number: S34001; MUID:93209281
A.Accession: S34003
A.Status: preliminary
A.Molecule type: mRNA
A.Residues: 1-113 <MARN
A.Residues: 1-113 <MARN
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C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;16-96/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-113 <MAR>
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Pred. No. 0.37;
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C;Comment: This is a Bence Jones protein.
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (C;Complex: An immunoglobulin to cases, such as IgA and IgM, the subunits associate into C;Superfamily: immunoglobulin region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin homology C;Keywords: immunoglobulin homology C;Meywords: 
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A;Title: Expression of a public idiotype by human monoclonal IgM directed to myelin-a A;Reference number: A61458; MUID:90039128
A;Accession: F61458
A;Status: preliminary
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Search completed:
Job time: 469 sec
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Best Local Similarity
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RESULT 2
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RESULT 3
KV4B_HUMAN
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SEQUENCE FROM N.A.
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MEDLINE-86041853; PubMed-2997712;
Klobeck H.G., Bornkamm G.W., Comb
                                                                         Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                        01-JAN-1988 (Rel. 06, Created)
01-JAN-1988 (Rel. 06, Last sequence upda
15-JUL-1999 (Rel. 38, Last annotation up
19 kappa chain V-IV region JI precursor.
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01-JAN-1988 (Rel. 05, Created)
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15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain V-IV region precursor (Fragment).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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P06313;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the European Bloinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
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HSSP; P01789; 1MCP.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
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01-APR-1988 (Rel. 07, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-IV region B17 precursor.
Homo sapiens (Human)
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                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                       MEDLINE-86041854; PubMed-2997713;
Marsh P., Mills F., Gould H.;
"Detection of a unique human V kappa IV germline cDNA probe.";
                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                Submitted (OCT-1986)
                                                                            Marsh
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                            myeloma MPC 11.";
Blochem, J. 171:337-347(1978).
Blochem, J. 171:337-347(1978).
-I- MISCELLANEOUS: THE MATURE CHAIN HAS 12 ADDITIONAL RESIDUES AT ITS AMINO END, DUE TO A TANDEM DUPLICATION OF 36 NUCLECTIDES AFTER THI CODON FOR RESIDUE 36. RESIDUE 42 CORRESPONDS TO THE AMINO-TERMINAL RESIDUE OF TYPICAL KAPPA CHAINS.
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SEQUENCE
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"Functional significance and evolutionary development
"functional regions of immunoglobulin variable-region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE OF 1-71 FROM N.A. MEDLINE=83001944; PubMed=6288267;
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Mammalia; Eutheria;
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MEDLINE=78186617; PubMed=418775;
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PIR; A01905; K4HU17.
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                                                                                                                                            "Sequence of the full-length immunoglobulin kappa-chain nveloma MPC 11.":
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-JUL-1999 (Rel. 38, Last annotation update)
kappa chain V-V region MPC11 precursor.
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Hoppe-Seyler's Z. Physiol. Chem. 351:1291-1295(1970).
-!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE
-!- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
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"The primary structure of a monoclonal kappa-type immunoglobulin !
"The primary structure of a monoclonal kappa-type immunoglobulin !
"The primary structure of a monoclonal kappa-type immunoglobulin !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
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RESULT 8

KU1_HUMAN STANDARD; PRT; 117 AA.

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DT 15_JUL-1996 (Rel. 01, Last sequence)
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OC Entaryota: Metazoa: Chordata; Craniata: Vertebrata: Butel
Mammalia: Entheria: Primates; Catarrhini: Hominidae: Homo
NCRI_TaxID-9606;
RN 11
Nature 200:9566; DubMed-6779204;
RN SEQUENCE FROM N.A.
RY MEDILINE-81129397; PubMed-6402305;
RN 1210
RP SEQUENCE FROM N.A.
RY MEDILINE-81129397; PubMed-6402305;
RN MEDILINE-31129397; PubMed-6402305;
RN MEDILINE-31129 MM; Pappa sequences have diverged recently duplicated human V kappa sequences have diverged by a convergion.

RI Later Politicated human V kappa sequences have diverged by a centities requires a license agreement (See http://www.isb
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FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
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COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
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P04326;
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                                                                                   Human immunodeficiency virus type 1 (clone 12) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus. NCBI_TaxID-11679;
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Arya S.K., Gallo R.C.; "Three novel genes of human T-lymphotropic virus type III: reactivity of their products with sera from acquired immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Klobeck H.G., Combriato G., Zachau H.G.; "Immunoglobulin genes of the kappa light clymphoid cell lines are closely related.";
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MEDLINE-85014148; PubMed-6091049;
                                           SEQUENCE FROM N.A. MEDLINE=86177573; PubMed=3008154;
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SMART; SM00406; IGV;
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                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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13-AUG-1987
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alizon M., Wain-Hobson S., Montagnier L., "Genetic variability of the AIDS virus: n of two isolates from African patients.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    deficiency syndrome patients.";

Proc. Natl. Acad. Sci. U.S.A. 83:2209-2213(1986).

-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus type 1 (MAL isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                             EMBL; X04415; CAA28015.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
MEDLINE=86245056; PubMed=2424612;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=11697;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; M11840; AAA44999.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRINTS; PR00055; HIVTATDOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                   1 46:63-74(1986).

FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE
                                                                                                                                                                                                                                                                                                            SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
                                                                                                                                                                                                                                                                                                                                       PROMOTER.
SUBUNIT: BINDS CYCLIN T1
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SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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M11840; TATSPCV12
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05, Last sequence update)
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Pred. No.
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RESULT 12

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21-JUL-1986
15-JUL-1999
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Mammalia; Eutheria; I
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Complete amino acid sequence of the variable domains of IgM anti-gamma globulins (Lay/Pom) with shared idiotypic specificities.";
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DOMAIN
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HSSP; P01607; 1REI.
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Capra J.D., Klapper D.G.;
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-JUL-1986 (Rel. 01, Last sequence up
-JUL-1999 (Rel. 38, Last annotation
kappa chain V-I region Lay.
                                                                                                                                                                                                                           Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nd. J. Immunol. 5:677-684(1976).

MISCELLANEOUS: THE SECOND AND THIRD HYPERVARIABLE REGIONS OF THIS CHAIN ARE IDENTICAL WITH THOSE OF THE HUMAN POM V-III KAPPA CHAIN, WITH WHICH IT SHARES CERTAIN IDIOTYPIC DETERMINANTS.

MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM AN IGM WITH ANTI-GAMMA GLOBULIN ACTIVITY.
                                                                                                                                             SGSGTDATAPASSLQAED 29
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13; Conservative
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BY SIMILARITY.
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21-JUL-1986 (Rel. 01, Creat
21-JUL-1986 (Rel. 01, Last
15-JUL-1999 (Rel. 38, Last
15 Last (Rel. 38) (Last)
                                                                                                                                                                                                                                                                                                                                                                 Epp O., Lattman E.E., Schiffer M., Huber R., Palm W.; "The molecular structure of a dimer composed of the variable portions of the Bence-Jones protein REI refined at 2.0-A resolution."; Biochemistry 14:4943-4952(1975).

-I. MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE INV (1,2)
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SEQUENCE
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DOMAIN
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SMART; SM00406; IGv; 1.
Tmmunoglobulin V region;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Epp O., Lattman E.E.,
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                                                                                                                                                                                                                                                                                                                                   PDB; 1REI;
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         39.68;
72.28;
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         Score 57;
Pred. No.
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COMPLEMENTARITY-DETERMINING-2.
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          0.49;
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                  Length 108;
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SGSGTDYTFTISSLQPED

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DE KVIC_HUMAN

LD KVIC_HUMAN

LD KVIC_HUMAN

AC P0159;

AC P0159;

DT 21-UUL-1986 (Rel. 01, Created)

DT 21-UUL-1986 (Rel. 01, Last sequence update)

DT 15-UUL-1999 (Rel. 38, Last annotation update)

DE 19 kappa chain V-I region BI.

OS Homo sapiens (Human)

OS Homo sapiens (Human)

"""heria; Primates; Catarrhini; Homi!
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ID KVIY_HUMAN

AC P80362;
DT 01-NOV-1995
DT 01-NOV-1995
DT 15-JUL-1999
DE 15 Kappa che
OC Homo sapiene
OC Structural
OC Homo structural
OC Homo structural
OC Homo sapiene
OC Homo
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01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
101-NOV-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation update)
17 kappa chain V-I region WAT.
Homo sapiens (Human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Huang D.-B., Chang C.-H., Ainsworth C., Bruenger A.T., Eulitz M., Solomon A., Stevens F.J., Schiffer M.;
"Comparison of crystal structures of two homologous proteins: structural origin of altered domain interactions in immunoglobulin light-chain dimers.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PDB; 1WTL; 01-NOV-94.
InterPro; JPR003006; Ig_WHG.
InterPro; IPR003596; Ig_V.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Characterization and preliminary crystallographic data on the VL-
related fragment of the human ki Bence Jones protein Wat.";
J. Wol. Biol. 147:185-193(1981).
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MEDLINE-81267384; PubMed-6167731;
Stevens F.J., Westholm F.A., Panagiotopoulos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE, AND X-RAY CRYSTALLOGRAPHY (1.9 MEDLINE-95086080; PubMed-7993911;
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FRAMEWORK-1,
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Pred. No.
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COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
BY SIMILARITY.
TN -> SD (IN REF. 2).
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                                                     Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Search completed: July 15, Job time: 1450 sec

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Braun M., Leibold W., Barnikol H.U., Hilschmann N.;
Braun M. eibold W., Barnikol H.U., Hilschmann N.;
"Principle of antibody structure. The primary structure of a
monoclonal kappa I-type immunoglobulin L-chain (Bence Jones protein
Bi). 3. The complete aminno acid sequence and the genetic
significance of the variability principles for the mechanism of
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HSSP; P01607; 1REI.
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-!- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE
-!- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antibody formation.";
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InterPro; IPR003596;
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COMPLEMENTARITY - DETERMINING - 3
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O40234 homo sapien
O9n749 homo sapien
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ALIGNMENTS

OPFTY9 PRELIMINARY; PRT; 399 AA.

Q9FTY9;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
PUTATIVE RECEPTOR SER/THR PROTEIN KINASE.

P0436E04.9.

Oryza sativa (Rice).

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RESULT

Q9FTY9

ID Q9

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DT Q9

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OT Q9

OT Q9

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         Query Match
Best Local Similarity
Matches 13; Conserv
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PRINTS; PR00109; TYRKINASE,
SMART; SM00220; S_TKC; 1.

SMART; SM00219; TYRC; 1.

PROSITE; PS00101; PROTEIN_KINASE_DOM; 1.

PROSITE; PS00109; PROTEIN_KINASE_TYR; 1.

ATP-binding; Kinase; Receptor; Transferase.

ATP-binding; Kinase; Receptor; Transferase.
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Sasakl T., Matsumoto T., Yamamoto K.;
Sasakl T., Matsumoto T., Yamamoto DNA, chromosome 1,
"Oryza sativa nipponbare(GA3) genomic DNA, chromosome 1,
clone:P0436E04.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases EMBL; AP002818; BAB16326.1; .
InterPro; IPR000719; Euk_pkinase.
InterPro; IPR002290; Ser_thr_pkinase.
InterPro; IPR001245; Tyr_pkinase.
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Submitted (FEB-1998) to the EMBL/GenBank/DDBJ databases.

CITYMICTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT A ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FRO PROMOTER (BY SIMILARITY).

R EMBL; AF649495; AAC68854.1; -.

R Interpro; IPR001831; HIV_Tat.

R Pfam; PF00539; Tat; 1.

R Pfam; PF00539; Tat; 1.

R PFAM; PF000559; HIVATDOMAIN.

R PFRINTS; PR00055; HIVATDOMAIN.

ACTIVATOR; Nuclear protein; RNA-binding; Transcription regulation activator; Nuclear protein; RNA-binding; Transcription regulations and the second 
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040231;
01-JAN-1998
01-JAN-1998
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071973;
01-AUG-1998
01-AUG-1998
01-DEC-2001
                                                                          STRAIN-HE93-A3TAT;
MEDLINE-97335179; PubMed-9191845;
Kirchhoff F., Greenough T.C., Hamacher M., Sulliv
Desrosiers R.C.;
"Activity of human immunodeficiency virus type 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=98440562; PubMed=9765443;
Mwaengo D.M., Novembre F.J.;
Molecular cloning and characterization
chimpanzees with pathogenic human immuno
infections.":
                                             and
                                                                                                                                                                                                                                                                                                                                                           Viruses; Retroid
NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Viruses; Retroid viruses; NCBI_TaxID=11676;
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disease progression
                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
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            tatl genes derived ase progression.";
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ATING REGULATORY PROTEIN).
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Last annotation updat
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                                                                                                                                                                 Sullivan J.L.
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                                             different rates
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Best Local S
Matches 15
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Best Local (
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                                                                                                                                                                                     O9NR59 PRELIMINARY; PRT; 1312 AA. 09NR59; 01-OCT-2000 (TrEMBLrel. 15, Created) 01-DEC-2000 (TrEMBLrel. 15, Last sequence update) 01-DEC-2001 (TrEMBLrel. 19, Last annotation update) SER/ARG-RICH PRE-MRNA SPLICING FACTOR SR-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases EMBL; AB002405; BAAZ4179.2; .
InterPro; IPR000515; BPD_transp.
PROSITE; PS00402; BPD_TRANSP_INN_MEMBR; UNKNOWN_1.
SEQUENCE 444 AA; 50427 MW; DD4053A374A00FA1 CRC64;
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TISSUE-LYMPHOID;
Abe Y., Takaoka Y.;
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PRINTS; PR00055; HIVTATDOMAIN.
Activator; Nuclear protein; RN
NON TER 71 71
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1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATION RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).

EMBL; AF000529; AAB62528.1; -.
EMBL; AF000529; AAB62528.1; -.
ENERGY IPRO1831; HIV_Tat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Primates;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUN-1998 (TREMBLRel. 06, 01-MAY-2000 (TREMBLRel. 13, 01-JUN-2001 (TREMBLREL. 17,
                                                                    Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
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01-JUN-1998
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                      EQUENCE
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hes 13; Conserv
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15; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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Kyriakopoulou
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 62; DB
Pred. No. 0.91
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 62; DB Pred. No. 0.14 0; Mismatches
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                                                                                             Craniata; Vertebrata; Catarrhini; Hominidae;
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
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Q9H7N4;
01-MAR-20Q1 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
01-DEC-2001 (TrEMBLrel. 19,
MEDIJINE-98440562; PubMed-9765443;
Mwaengo D.M., Novembre F.J.;
"Molecular cloning and characterization of viruses chimpanzees with pathogenic human immunodeficiency
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Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF254411; AAF87552.1; ...
InterPro; IPR00295; P.rich_extensn.
PRINTS; PR01217; PRICHEXTENSN.
SEQUENCE 1312 AA; 139269 MW; OCB1C87C963C52BD CRC64;
                                                                                                                                                                                                                                                              Human immunodeficiency virus type Viruses; Retroid viruses; Retrovir
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InterPro; IPR002965; P_rich_extensn
PRINTS; PR01217; PRICHEXTENSN.
                                                                                                                                     SEQUENCE FROM N.A.
STRAIN-HIV-1JC;
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A WAGNIG D.M., NOVEMBRE F.J.;
SUBMILTED (FEB-1998) to the EMBL/GenBank/DDBJ databases.
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2 -|- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDII
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R EMBL; AF049494; AAC68845.1; -.
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SEQUENCE 101 AA; 11552 MW; DELEFFCC34D9CDB8 CRC64;
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Carr J.K., Avila M., Gomez-Carrillo M., Salomon H., Hic Carr J.K., Avila M., Gomez-Carrillo M., Russell K.L., Watanaveradej V., Pando M., Negrete M., Russell K.L., Birx D.L., Andrade R., Vinoles J., McCutchan F.E.; "Diverse BF Recombinants have spread widely since the intro South America.";
HIV-1 into South America.";
ALDS 0:0-0(2001).
EMBL, AY037277; AAL12713.1; -.
Viruses;
                        Human immunodeficiency virus
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Viruses; Retroid viruses; Retroviridae;
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J. Virol. 72:8976-8987(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YGRKKRRORRRPRQGSQAHQASLSEQPTSQLRGD
                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
  Retroid viruses;
                                                                 5 (Tremblrel 01, C:
5 (Tremblrel 01, L:
1 (Tremblrel 19, L:
N (TRANSACTIVATING)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (TrEMBLrel. 1 (TrEMBLrel. 1 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  101
                                                                                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAL12713.1; -.
AA; 11572 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                   42.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42.48;
44.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19,
19,
rus type 1.
Retroviridae;
                                                                                                                                                                                                                                                                                                                                                                                           ; Score 60.5; DB; Pred. No. 0.33; 1; Mismatches
                                                                 Created)
Last sequence update)
Last annotation update)
S REGULATORY PROTEIN) (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                 ------DATAPASS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Last sequence up
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ω,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 61; DB
Pred. No. 0.28
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13A12A179D2EF930 CRC64;
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                                                                                                                                                                                      72
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  Lentivirus.
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                                                                                                                                                                                   A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B
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                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       update)
                                                                                                                                                                                                                                                                                                                                                                                        7;
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                                                                                                                                                                                                                                                                                                                                                 24
                                                                                                                                                                                                                                                                                                      87
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BINDING TO THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         introduction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sanchez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  regulation
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                                                                                                                                                                                                                                                                                                                                                                                                                                              101;
                                                                                                                                                                                                                                                                                                                                                                                               17;
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                                      Query Match
Best Local S
Matches 11
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Best Local S
Matches 14
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J. Virol. 69:3273-3281(1995).

-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                                                                         NON_TER
SEQUENCE
                                                                                                                    01-MAY-1997 (TrEMBLrel. 03, Created)
01-MAY-1997 (TrEMBLrel. 03, Equence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN) (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                         P90117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=95264414; PubMed=7745674;
Diaz R.S., Sabino E.C., Mayer A., I
"Dual human immunodeficiency virus
                                                                                                                                                                                                                                                                       Human immunodeficiency virus type 1. Viruses; Retroid viruses; Retroviridae; NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Activator; Nuclear protein; NON_TER 72 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Dual human immunodeficiency virus type 1 infection and recin a dually exposed transfusion recipient. The Transfusion
                                                                                        Activator; Nuclear protein; RNA-binding; Transcription regulation NON_TER 72 72
SEQUENCE 72 AA; 8214 MW; F79D7B8D0205EEB0 CRC64;
                                                                                                                                                                                                                                                      SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PRO0055; HIVTATDOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID-11676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                      47
 47
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J, Ull185; AAA78871.1; -

arPro, IPR001831; HIV_Tat.
                                                                                                                                                                                                                                                                                                                                                                                                                                        YERKKRRORRRSGSGTDATAPASSLO
                   YERKKRRQRRRSGSGTD
                                                                                                                                                                                                                                                                                                                                                                                                                     YGRKKRRORRRAPQGNQAHQVSLSKQ
 YGRKKRRQRRRASQGSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PF00539; Tat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 53.8
14; Conservative
                                                 Similarity
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                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
                                                41.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41.78;
 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA-binding; Transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score
Pred
                                                Score 60;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A., Mosley J.W.,
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                                                                                                                                                                                                                                                                                                                                                                                                                       72
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                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                    72
                                                   0
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0.27;
                                                           DB
                                                                                                                                                                                                                                                                                    Lentivirus
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                                                           Length
                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         recombination
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                                                                                                                                                                        THE LTR
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Best Local S
Matches 14
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P90128;
01-MAY-1997
01-MAY-1997
01-DEC-2001
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Viruses; Retroid viruses; Retroviridae;
NCBI_TaxID=11676;
[1]
                                                                                                                                                                                                                                                             Q90MK8;
Carr J.K., Avila M., Gomez-Carrillo M., Salomon H., Hierholzer Watanaveeradej V., Pando M., Negrete M., Russell K., Sanchez J., Birx D., Andrade R., Vinoles J., McCutchan F.; Submitted (MAY-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF385936; AAK69333.1; .

SEQUENCE 101 AA; 11416 MW; 360605029A87A327 CRC64;
                                                                                        Carr J.K., Avila M., Gomez-Carrillo M., Salomon H., Hi Watanaveeradej V., Pando M., Negrete M., Russell K., S Birx D., Andrade R., Vinoles J., McCutchan F.; "Diverse BF recombinants have spread widely since the HIV-1 into South America.";
                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PR00055; HIVTATDOMAIN. Activator; Nuclear protein; RN NON_TER 72 72
                                                                                                                                                                                          Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
                                                                                                                                                                                                              TAT
                                                                                                                                                                                                                               01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                      pfam; PF00539; Tat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                          STRAIN-ARMA159;
                                                                                                                                                    STRAIN-ARMA159;
                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                NCBI_TaxID=11676;
                                                                                                                                                                                                                         TAT PROTEIN
                                                                       SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                         47 YGRKKRRQRRRSPQGSQTHQVSLSKQ 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11
                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                                           1 YERKKRRORRRSGSGTDATAPASSLQ 26
                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TrembLrel. 03, C: (TrembLrel. 03, L: (TrembLrel. 19, L: (TrembLrel. 19, L: (Transactivating)
                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                     N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                8337 MW;
                                                                                                                                                                                                                                                                                                                                                                         41.7%;
53.8%;
                                                                                                                                                                                                                                  19,
19,
                                                                                                                                                                                                                               Created)
Last sequence Last anno
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Last annotation update)
GREGULATORY PROTEIN) (FRAGMENT).
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                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA-binding; Transcription regulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Created)
                                                                                                                                                                                                                                                                                                                                                                                                                4D0AED1145E6645C CRC64;
                                                                                                                                                                                                                                                                                                                                                                        Score 60; I
                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                 sequence up
                                                                                                                                                                                                                                                                          101
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                                                                                                                                                                                            Lentivirus
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                                                                                                                                          Hierholzer
                                                                                                                                  Sanchez
                                                                                                              introduction
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Best Local Similarity
Matches 13; Conser
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01-OCT-2000 (TrEMBLrel. 15
01-OCT-2000 (TrEMBLrel. 15
01-DEC-2001 (TrEMBLrel. 16
01-DEC-7001 (TrEMBLrel. 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local :
               Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscu
Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID-7227;
                                                                                                   01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 01-JUN-2001 (TrEMBLrel. 17, CG6184 PROTEIN.
                                                                                                                                                                                      Q9VDV0;
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Interpro; IPR001752; kinesin.
Pfam; PF00225; kinesin; 1.
PRINTS; PR00380; KIMESINHEAVY.
SMART; SM00129; KISc; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS00411; KINESIN_MOTOR_DOMAIN1; 1.
PROSITE; PS50067; KINESIN_MOTOR_DOMAIN2; 1.
ATP-binding; Coiled coil; Microtubules; Motor protein.
NON TER 820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "A physical map of the Leishmania Genome Res. 8:135-145(1998). EMBL; AL157416; CAB75648.1; -. HSSP; P17119; 3KAR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ivens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murphy L., Harris D., Barrell B.G.; Submitted (FEB-2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ivens A.C., Lewis S.M.,
Smith D.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-FRIEDLIN;
MEDLINE-98146435; PubMed-9477341;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
STRAIN-FRIEDLIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leishmania major.
Eukaryota; Euglenozoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                      294
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
tes 13; Conserv
                                                                                                                                                                                                                                                                                                                     2
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                                                                                                                                                                                                                                                                                                             ERKKRRORRRSGSGTDATAPASSLQAE 28
                                                                                                                                                                                                                                                                                   EQRRLRQRRRGGKRADATAVMMQAQQE 320
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                                                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                   AA;
                                                                                                                                                                                                                                                                                                                                                                41.7%;
48.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                  92423 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               41.7%;
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15, Last sequence update)
19, Last annotation update)
(FRAGMENT).
                                                                                                                             Created)
Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                 Score 60; DB
Pred. No. 3.3;
3; Mismatches
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                                                                                                                                                                                                    PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  31B286D811B7D648 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  major
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                                                                                                                                                                                                                                                                                                                                                                                DB 5;
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RA Abril J.F. Abjayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ra Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Ra Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Ra Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Ra Burris K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Ra Burris K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Ra Borkova D., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M., Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Ra Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., Ra Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Ra Glodek A., Gong F., Gorrell J.H., Gu Z., Gener S., Fleischmann W., Ra Harris N.L., Harvey D., Helman T.J., Hernandez J.R., Houck J., Ra Hostin D., Houston K.A., Howland T.J., Hernandez J.R., Houck J., Ra Hostin D., Houston K.A., Howland T.J., Hernandez J.R., Houck J., Ra Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Lin X., Li J., Li Z., Liang Y., Lin X., Ra Liu X., Mattel B., McIntosh T.C., McLeodd M.P., McPherson D.L., Ra Kill M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., Nixon K., Nusskern D.R., Pacleb J.M., Palazolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G., Ra Meisen D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., Ra Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G., Ra Ra Maris S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ra Alliu X., Moodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ra Alliu X., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ra Zheng X.H., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ra Zheng X.H., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Pittman G.S., Pan M., Zhang G., Zhao Q., Zheng L., Ra Maris M.R., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Pittman G.S., Polin G.M., Venter J.C., Ra Gibls R., Wyers E.M., Ra Kandra M., Ra
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P90148
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-1997
01-MAY-1997
01-DEC-2001
SEQUENCE FROM N.A.
STRAIN-PATIENT LTS5;
MEDLINE-99049627; PubMed-9833881;
                                                                                                                                                                                                                                                Human immunodeficiency virus type 1.
Viruses: Retroid viruses; Retroviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
Basilow B. Basilow B. Bayendala. J. Ba
                                                                                                                                                                                                             NCBI_TaxID=11676;
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7 (TremBLrel. 03, La
1 (TremBLrel. 19, La
N (TRANSACTIVATING )
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46.48;
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Last annotation update)
G REGULATORY PROTEIN) (FRAGMENT).
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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         protein search, using sw model
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14.104 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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144
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                                                                                                                                                                                                                       Length
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Interleukin-5 huma
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6	9 6	3 1	8	18	18	21	20	20			21	20	20	20	20	16	16	17	22	21	20	22	15	18	18	15	14	22	21	21	21	21	21	21	
KK134311	22424211	ALCVENA	AAW10545	AAW10544	AAW10547	AAB03714	AAY50690	AAY05264	AAW21934	AAP80894	AAB07966	AAY18118	AAY18126	AAY23771	AAY23779	AAR70202	AAR75355	AAW06443	AAB78867	AAY77598	AAY57181	AAG65565	AAR61240	AAW27546	AAW15537	AAR45605	AAR30144	AAB62769	AAB18881	AAB18879	AAB18873	AAB18871	AAB18869	AAB18863	
igm ancibody CEM i	3	10146CH 11111	murine	murine.	Wild type murine a	1110	n4 VL C	24-31	ligh	region	ht chai	chain	chain		_	antibod	Humanized antibody	egion.	human Flt-1	human	acid	acid	Ţ,	light c	Anti-TGF beta-1 sc		AH variable	HIV-1	acid	acid		acid	acid	Amino acid sequenc	

ALIGNMENTS

AAW62807 RESULT

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AAW62807 standard;

Peptide;

82 ₿

23-SEP-1998 AAW62807;

(first

entry)

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Green L,
Mendez M;
 New transgenic non-human mammals
                     WPI; 1998-333314/29.
                                                                                                                                                                                     Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa; production; antibody.
                                                                                       03-DEC-1996;
                                                                                                           03-DEC-1997;
                                                                                                                              11-JUN-1998.
                                                                                                                                                 W09824893-A2
                                                                                                                                                                     Homo sapiens.
                                                                    (ABGE-) ABGENIX INC.
                                                                                                                                                                                                                                              Amino acid sequence of a human antibody fragment.
                                                 Jakobovits
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an inactivated
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iccus, so that the mammal does not display normal becar to the modified genome also has an inserted human heavy chain Ig locus in germline configuration, the human heavy chain Ig locus comprising a human configuration, the human heavy chain Ig locus comprising a human ICC genes, human D-H genes, and human V-H genes and an inserted human N-H genes, human Ig locus in germline configuration, the human kappa light chain Ig locus comprising a human kappa constant region, J-kappa genes, and V-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The CC transgenic animals have a near complete human Ig locus, including both a human heavy chain locus and a human kappa light chain locus. They can be used for the production of human happa light chain locus. They can the mice will produce antibodies to IL-8, EGFR or TNF- alpha
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                                                                                                                                                                                                                                                                                                                                      kappa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specification describes a transgenic non-human mammal which has genome modifications that comprise an inactivated endogenous immunoglobulin (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW62793-822 represent fragments of human antibodies produced by transgenic Xenomice, created using the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunoglobulin locus and a near complete human immunoglobulin locus, used for production of human antibodies
 New transgenic non-human mammals - having an inactivated immunoglobulin locus and a near complete human immunogloused for production of human antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW62808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW62808 standard; Peptide; 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                      03-DEC-1996;
                                                                                                                                                                                                   03-DEC-1997;
                                                                                                                                                                                                                                                               W09824893-A2
                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                   Amino
                                                                                                                                       (ABGE-) ABGENIX INC
                                                                                                                                                                                                                                                                                                                                    Human; immunoglobulin; Ig; transgenic; non-human mammal;
inactivated endogenous Ig locus; B-cdi development;
human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene;
kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SGSGTDATAPASSLQAED 29
                                                              1998-333314/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sgsgtdftltisslqaed
                                                                                                                                                                                                                                                                                                                                                                                                                   acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                         Jakobovits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         82 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                      9605-0759620
                                                                                                                                                                                                    97WO-US23091.
                                                                                                                                                                                                                                                                                                                                                                                                                   of a human antibody fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43.8%;
                                                                                                           ۶,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . 88;
                                                                                                           Klapholz S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 63; DB Pred. No. 0.2; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A
                                                                                                           Kucherlapati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 19;
0.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      normal B-cell development. The heavy chain Ia looms in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
                 immunoglobulin locus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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constitution describes a transgenic non-human mammal which has genome comodification that comprise an inactivated endogenous immunoglobulin (Ig) cous, so that the mammal does not display normal B-cell development. The comodified genome also has an inserted human heavy chain Ig locus in the configuration, the human heavy chain Ig locus comprising a human constant region and regulatory and switch sequences, human J-H cous in germline configuration, the human heavy chain Ig locus comprising a human configuration, the human kappa and an inserted human kappa iight constant Ig locus in germline configuration, the human kappa light constant Ig locus comprising a human kappa constant region, J-kappa genes, and human kappa constant region, J-kappa genes, cand V-kappa genes, where the number of V-H and V-kappa genes inserted constant region, J-kappa genes, and human kappa constant region, J-kappa genes, cand V-kappa genes inserted constant region, J-kappa genes, where the number of V-H and V-kappa genes inserted consuments and including both a human happa light chain locus. They can be used for the production of human happa light chain locus. They can be used for the production of human antibodies when exposed to particular antigens e.g. when exposed to human IL-8, EGFR or TNF- alpha constrively.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                δã
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                 Antibody 24-31; variable light chain; humanised antibody; gene therapy; CD40 binding inhibitor; gp39; autoimmune disease; food-related allergy; graft-versus-host disease; graft rejection; intestinal inflammation; reversible obstructive airways disease; allergy; transplant rejection; asthma; leukaemia; lymphoma; immunosuppressant; cellular therapy;
                                                                                                                                                                                                                                                                                                            Homo sapiens.
Synthetic.
 Claim 7; Page 84; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY05269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY05269 standard; Protein; 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 79; 128pp; English.
                                    New humanized
                                                                                                     Black A,
                                                                                                                                                                          08-SEP-1997;
                                                                                                                                                                                                            08-SEP-1998;
                                                                                                                                                                                                                                             18-MAR-1999.
                                                                                                                                                                                                                                                                             WO9912566-A1
                                                                                                                                                                                                                                                                                                                                                Mus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antibody
                                                                    WPI; 1999-229142/19
                                                                                                                                       (IDEC-) IDEC
                                                                                                                                                                                                                                                                                                                                                                                   .mmune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 SGSGTDATAPASSLQAED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               52 sgsgtdftltisslqaed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-31 humanised VL chain sequence
                                                                                                       Hanna N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  represent fragments of human antibodies produced by enomice, created using the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                         PHARM CORP
                                    antibodies to human
                                                                                                                                                                        97us-0925339
                                                                                                                                                                                                            98WO-US18163
                                                                                                                                                                                                                                                                                                                                                                                   inhibitor;
                                                                                                        Newman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  43.8%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29
                                                                                                                                                                                                                                                                                                                                                                                    cancer
                                                                                                        RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 63; DB 19; Pred. No. 0.24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                        Padlan
                                                                                                        EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Key
Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibody; humanised; murine; human; framework region; complementarity defined to the state of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR52059 standard; Protein; 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modelling;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Light chain variable region of human KV4B antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-OCT-1996
     09-SEP-1992;
                                                         07-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                         93EP-0307051.
                                                                                                                                                                                                                                                   /note=
96..104
                                                                                                                                                                                                                                                                                                                                                                /note=
56..62
                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
41..55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note-
24..40
                                                                                                                                                                                                                                                                                                                                   /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                             /label= complementarity_determining_region_l
/note= "CDR 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
1..23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= framework_region_1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complementarity determining region; reshaping; residue; modify.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43.8%;
77.8%;
                                                                                                                                                                                                                                                                             "FR 3"
                                                                                                                                                                                                                                                                                                                                   "CDR 2"
                                                                                                                                                                                                                                                                                                                                                                                         "FR 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "FR 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 63;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             heavy chain; light; variable;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 20;
0.27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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RESULT
AAW27527
ID AAW2
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Best Local Similarity
Matches 14; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modification of a rodent antibody (Ab) or fragment by resurfacing in order to produce a humanised rodent Ab can be determined by calculating homology between murine and human Ab antibody surfaces. In order to test the resurfacing approach of the invention, three humanisation experiments were set up: (1) traditional loop grafting; (2) resurfacing approach using most similar chain; and (3) resurfacing approach using most similar chain; and (3) resurfacing approach using murine anti-N901 Ab (see AAR52055). Experiment 3 was carried out using the present sequence which represents the human KV4B Ab light chain variable region with 71 percent homology with anti-N901 Ab. N901/KV4B (AAR52060) was prepd. by resurfacing.
                                                                           (ASAK ) ASAHI I
(NIKK-) NIKKA I
(TORI ) TORII :
(TSUR/) TSURA
                                                                                                                                                                                                                                                                                             high affir IgE; MAb;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Method of resurfacing of rodent antibodies to produce humanised antibody forms - for producing non-human antibodies with improved therapeutic efficiency by presenting human surface on V-region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Fig 4A; 230pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1994-120230/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Guild BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PEDE/) PEDERSEN J T. (IMMU-) IMMUNOGEN INC
                                                                                                                                                                                                   29-JUL-1997
                                                                                                                                                                                                                            JP09191886-A
                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                 chimeric; treatment;
                                                                                                                                                                                                                                                                                                                     Complementarity
                                                                                                                                                                                                                                                                                                                                                 Light chain variable region
                                                                                                                                                                                                                                                                                                                                                                           16-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                       AAW27527;
                                                                                                                                                                                                                                                                                                                                                                                                                                AAW27527 standard; Protein; 112 AA
                                                   WPI; 1997-429186/40.
                                                                                                                                              19-JAN-1996;
                                                                                                                                                                        19-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                                                                                                                                                         տ
                                                                                                                                                                                                                                                                                            lementarity determining region; CDR; murine; mouse; human;
affinity; immunoglobulin E; receptor; monoclonal antibody;
MAb; light chain; variable region; humanised; semi-chimeri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sgsgtdftltisslqaed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SGSGTDATAPASSLQAED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    111 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                        WHISKEY
                                                                                                                   BREWERIES LTD
                                                                                         YAKUHIN
                                                                                                                                              96JP-0024816
                                                                                                                                                                        96JP-0024816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43.8%;
                                                                                                                                                                                                                                                                               prevention;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rees
                                                                                                                                                                                                                                                                                                                                                  of human CRA2 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 63; D. Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                 disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roguska MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1
0.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15; Length 111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.
                                                                                                                                                                                                                                                                                             semi-chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Humanised, semi-chimeric and chimeric antibodies against human high-affinity IgE receptor – useful medicinally and have low antigenicity in humans

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                New engineered antibodies and Plasmodium infection - contg. corresp. nucleic acid, vectors
Naturally-occurring
                                                                                                                                                                                                                                          Charoenvit
Sadoff JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence, the light chain variable region of the human antibody (Ab) CRA2, was used in the preparation of a humanised or semi-chimeric monoclonal Ab (MAb), comprising complementarity determining regions (CDR) from a menti-human high affinity immunoglobulin E (IgE) receptor, MAb. The humanised, semi-chimeric or chimeric MAb can be used to treat or prevent diseases, specifically allergies, associated with the receptor, and has very
                                                                                                                                                                                                                                                                                                                                                                                                                                   08-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            complementarity determining region; fusion protein; murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                                                             (SMIK
                                                                                                                                                                                                                                                                                                                                                                                         09-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9405690-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR50313;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR50313 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Monoclonal antibody; Plasmodium falciparum; CDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Humanised light chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                         (USNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6
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                                                                                                                                                                                             1994-101115/12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14;
                                                                                                                                                                                                                                                                                                    SMITHKLINE BEECHAM
US SEC OF ARMY.
US SEC OF NAVY.
                                                                                                                                                                         AAQ44827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 14; 26pp;
                                                                                                                                                                                                                                      Y, Hoffman S, Sylvester DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    112 AA;
                                          ຸນ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                        98pp;
                                                                                                                                                                                                                                                                                                                                                                                         92US-0941654
                                                                                                                                                                                                                                                                                                                                                                                                                                 93WO-US08435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= CDR2
95..103
/label= CDR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers 24..40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= CDR1
56..62
(AAQ44841-42) and
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77.8%;
                                          English.
                                                                                    vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86
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Gross
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Pred. No.
                                                                                fusion proteins murine antibody s and transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              region Pfhzlc1-1.
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synthetic
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0.28;
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Y CDR sequences,
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(AAQ44825-28)
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Best Local
                 Naturally-occurring (AAQ44841-42) and synthetic (AAQ44825-28) variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb NFS2, its variable chain peptides, CDRs, functional fragments, Fab fragments, and analogs are useful in prodn. of fusion
                                                                                                                                                                                                                                                                                                                                       Charoenvit Y, Hoffmar Sadoff JC, Sylvester
                                                                                                                                                                                                                                                                                                                                                                                                                           (SMIK )
                                                                                                                                                                                       New engineered antibodies and fusion proteins Plasmodium infection - contg. murine antibody corresp. nucleic acid, vectors and transformed
                                                                                                                                                  Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibody; Plasmodium falciparum; CDR; complementarity determining region; fusion prote murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                            (USNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb its variable chain peptides, CDRs, functional fragments, Fab fragments, and analogs are useful in prodn of fusion proteins, esp. engineered antibodies. These prods are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      used to protect humans against Plasmodium infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        71
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14; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                          OF ARMY.
                                                                                                                                                                                                                                                                                                                                                                  Hoffman
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                                                                                                                                           98pp;
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24..40
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77.8%;
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                                                                                                                                               English
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Gross
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                              S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    functional fragments,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fusion protein;
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                                                                                                                                                                                       s for preventing Y CDR sequences, ad cells
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Best Local
                                                                                                                                                                                                                                                                     06-JUN-1995;
23-DEC-1994;
06-JUN-1995;
                                                                                                                                                                                                           (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antibody engineering; humanised antibody; chimeric antibody; Fab; interleukin-5; IL-5; eosinophil; asthma; allergic rhinitis; atopic dermatitis; therapy; diagnosis; light chain; VL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         used
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                New monoclonal antibody to human interleukin-5 - used to produce products for the treatment and diagnosis of conditions associated with excess eosinophil prodn., e.g asthma etc.
                                                                                                                                                                                                                                                                                                                                                                                        11-JUL-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Humanised 2B6 antibody light chain variable region.
                                                                                                                                                        Holmes
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                                                                                              1996-333976/33.
DB; AAT34094.
                                                                                                                                                    SD,
                                                                                                                                                                                                             SMITHKLINE SMITHKLINE
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94US-0363131.
95US-0467420.
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104..113
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Theisen TW;
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                                                                                                                                                                               Region
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Region
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          21-JUN-1996;
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                              97WO-US10769
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77.8%;
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Pred. No. 0.
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RESULT 10
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KW P5
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Best Local Similarity
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Polynúcleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
                                                                                                     N-PSDB; AAA96134.
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rheumatoid arthritis; coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amino acid sequence of anti-p53 antibody light chain clone 163.2.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                113 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                               99AU-0009321
                                                                                                                                                                                                                                                                                                                                                                                                                immune response; vaccine; gene therapy; cancer;
hritis; coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein; 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43.8%;
77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No.
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The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies

Claim 30; Page 136; 163pp; English.

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Matches
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Best Local
The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in
                                                                                                                                                                                                                                                                              Ward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, head and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, gastric cancer, brain cancer, bladder cancer, prostate cancer and urinary/genital tract cancer, oesophageal cancer, mesenchymal tumours, e.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides
                                                                                                                        Claim
                                                                                                                                                                 Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
                                                                                                                                                                                                                              N-PSDB; AAA96140.
                                                                                                                                                                                                                                                                                                                                                 19-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amino acid sequence of anti-p53 antibody light chain clone 163.7.
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                                                                                                                                                                                                                                                                                                               (SVIN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rheumatoid arthritis;
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                                                                                                                                                                                                                                            2000-638249/61.
                                                                                                                         30;
                                                                                                                                                                                                                                                                                                           ST VINCENT'S HOSPITAL SYDNEY LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                      Page
                                                                                                                                                        and coronary heart disease
                                                                                                                                                                                                                                                                              Coomber DWJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in pharmaceutical compositions, which additionally contain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                        138;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Arg encoded by CCA"
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                                                                                                                      163pp;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vaccine; gene therapy; cancer;
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RESULT 3
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Best Local Similarity
        The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, head and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, gastric cancer, brain cancer, bladder cancer, prostate cancer and urinary/genital tract cancer, esophageal cancer, mesenchymal tumours,
                                                                                                                                                                                                                                                                      Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease -
                                                                                                                                                                                                                                                                                                                                                                                              Ward
                                                                                                                                                                                                                                    Claim 30; Page 144; 163pp; English.
                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAA96146
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77.8%;
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Pred. No.
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of epithelial origin,

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RESULT I AAB18871 ID AAB18871 ID AAB18871 AC AAB1871 AC
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Best Local :
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rheumatoid arthritis;
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                                                   The present sequence represents the light chain of an antibody reactive capainst p53. The antibody is obtained from a vertebrate host expressing can immune response against a naturally occurring disease. The antibodies care useful in pharmaceutical compositions, which additionally contain chalacters, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a covertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as compared to the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, head and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, castric cancer, brain cancer, bladder cancer, prostate cancer and curinary/genital tract cancer, oesophageal cancer, mesenchymal tumours, cover and cancer, sarrooma, and hemopoietic tumours, e.g. B cell lymphoma.
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rheumatoid arthritis; coronary heart disease.
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   July 15, 2002, 12:59:20; Search time 75.67 Seconds (without alignments) 9.361 Million cell updates/sec
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US-08-54-840-13
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US-08-525-539A-81
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US-08-467-420A-21
US-08-467-769A-21
US-08-677-769A-21
US-08-677-769A-21
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US-08-677-769A-21
US-08-677-769A-21
US-08-525-539A-80
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Query Match
Best Local Similarity 77.
Matches 14; Conservative

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ALIGNMENTS RESULT 1 US-08-54-840-13	28 63 43.8 133 3 US-08-463-903-4 Sequence 4, Appli 30 63 43.8 135 3 US-08-812-586-46 Sequence 46, Appli 31 63 43.8 135 3 US-08-828-741B-11 Sequence 11, Appli 32 63 43.8 155 4 US-08-828-741B-11 Sequence 20, Appli 33 63 43.8 171 3 US-08-463-903-20 Sequence 20, Appli 35 63 43.8 241 2 US-07-935-695-20 Sequence 20, Appli 36 63 43.8 260 3 US-08-463-903-2 Sequence 26, Appli 37 63 43.8 260 3 US-08-463-903-2 Sequence 27, Appli 38 63 43.8 275 3 US-08-463-903-2 Sequence 27, Appli 39 63 43.8 275 3 US-08-463-903-2 Sequence 6, Appli 63 43.8 275 3 US-08-463-903-2 Sequence 22, Appli 63 43.8 275 3 US-08-463-903-2 Sequence 22, Appli 63 43.8 285 3 US-08-463-903-2 Sequence 22, Appli 63 43.8 342 3 US-08-828-741B-6 Sequence 22, Appli 63 43.8 342 3 US-08-828-741B-6 Sequence 6, Appli 64 43 63 43.8 342 3 US-08-828-741B-6 Sequence 6, Appli 64 65 43.8 342 3 US-08-828-741B-6 Sequence 6, Appli 65 63 43.8 342 4 US-09-160-567-6 Sequence 4, Appli 65 65 65 65 65 65 65 65 65 65 65 65 65
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                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (703) 836-2021
[INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 80 amino acids
                                                  APPLICANT: NEWAAN, ROLand A.
TITLE OF INVENTION: HUMANIZED ANTIBODIES TO HUMAN 9D39,
TITLE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPE
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/554,840 FILING DATE: 07-NOV-1995 CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REPERENCE/DOCKET NUMBER: 012712-127
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 336-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: HUMANIZED ANTIBODIES TO HUMAN 9p39,
TITLE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPEUTIC USE THEREOF
                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acids
                                                                                                                                                                                                                                                                                                          47 SGSGTDFTLTISSLOAED 64
                                                                                                                                                                                                                                                                                                                                            12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 22313-1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     47 SGSGTDFTLTISSLOAED 64
                  ADDRESSEE:
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P.O. Box 1404
 P.O.
                                                                                                                       BLACK, Amelia
HANNA, Nabil
PADLAN, Eduardo A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NEWMAN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HANNA, Nabil
                                                                                                                                                                                                                                                                                                                                                                                Conservative
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Burns, Doane, Swecker & Mathis
O. Box 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roland
                                                                                                                                                                                                                                                                                                                                                                                                43.8%;
77.8%;
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Pred. No. (
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                                                                     AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                4.
                                                                                                                                                                                                                                                                                                                                                                                                               Length 80;
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US-08-525-539A-81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 81, Application US/08525539A Patent No. 6309636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                              ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, VG

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPAX: (703) 836-2021 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,612
REFERRNCE/DOCKET NUMBER: 276:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: DO COUTO,
APPLICANT: CERTANI,
APPLICANT: PETERSONI,
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESSE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08 FILING DATE: 07-NOV-1995 CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Rela
                                                                                                         APPLICATION NUMBER: US/08/525,539A FILING DATE: 14-SEP-1995 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     65 SGSGTDFTLTISSLQAED 82
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TOPOLOGY: li
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STATE: Virginia
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ZIP: 22313-1404
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                                                                                                                                                                                                                                                                                                               Palo Alto
:: CA
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Similarity 77.8%;
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NVENTION: RECOMBINANT PEPTIDES DERIVED FROM THE
NVENTION: MC3 ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DO COUTO, FERNANDO J.R. CERIANI, ROBERTO L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         107 amino acids
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                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                                                                                                                                              METHODS OF HUMANIZING ANTIBODY PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Release #1.0, Version #1.30
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                                        27633-20001.21
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Pred. No. 0
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                                                                                                                                                                                     Version #1.30
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RESULT 6
US-08-467-420A-21
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; MOLECULE TYPE:
US-07-942-245-30
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MEDIUM TYPE: Floppy disk
COMPUTER: HP 9000/700 Workstation
OPERATING SYSTEM: UNIX
SOFTWARE: IN house
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/07/942,245
FILING DATE: 09-SEP-1992
CLASSIFICATION: 530
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
TELEFAX: (202) 293-7860
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 5
US-07-942-245-30
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                                                                                                       Matches
                                                                                                                 Query Match
Best Local Similarity
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Best Local Similarity 77.8
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TELEX: 706141
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: GUILD,
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                           TYPE: amino acid
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CITY: Washington
STATE: D.C.
                                                    71
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                                                                          12 SGSGTDATAPASSLQAED 29
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STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                   SGSGTDFTLTISSLQAED 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/07942245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Sughrue, Mion, Zinn, Macpeak & Seas 2100 Pensylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REES,
ROGUSKA,
                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     United States
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                                                                                                                                                                                                linear
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Anthony R.
Anthony R.
Michael A.
Braydon C.
SURFACE RESIDUE VENEERING O
                                                                                                                 43.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTIBODIES
522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43.8%;
77.8%;
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                                                                                                                 Score 63; DB 1;
Pred. No. 0.043;
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Pred. No. 0.041;
                                                                                                      Mismatches
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                                                                                                                             Length 112
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: Sequence 21, Application US/08470110A

: Patent No. 5693323
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                                                                     GENERAL INFORMATION:
APPLICANT: Ames, R
APPLICANT: Appelba:
APPLICANT: Chaiken
APPLICANT: Cook, R
                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21, Application US/08467420A Patent No. 5683892
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 610 270-5090 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
TOPOLOGY: line
MOLECULE TYPE: F
                    APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 950282
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
TELEPAX: 610 270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: SmithKline Beecham Corp./Corporate ADDRESSEE: Intellectual Property STREET: P. O. Box 1539-UW2220 CITY: King of Prussia STATE: Pennsylvania COUNTRY: USA ZIP: 19406-0939
                                                                                                                                                                                                                                                                       71 SGSGTDFTLTISSLQAED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE:
                                                                                                                                                                                                                                                                                                        12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                              Local Similarity es 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : 113 amino acids
amino acid
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Appelbaum, Edward
Chaiken, Irwin M.
                                 Cook, Richard M. Gross, Mitchell S. Holmes, Stephen D.
                                                                                        Appelbaum, Edward Chaiken, Irwin M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gross, Mitchell S.
Holmes, Stephen D.
 McMillan, Lynette J
Theisen, Timothy W.
                                                                                                                            Ames, Robert S.
                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                            Score 63; DB 1; Pred. No. 0.043; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                              Length 113;
                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                            Gaps
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CORRESPONDENCE ADDRESS: TITLE OF INVENTION:

NUMBER OF SEQUENCES:

Recombinant IL5 Antagonists Useful in Treatment of IL5 Mediated Disorders

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Best Local (
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INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
                                                                                APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful :
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp./Corporate
STREET: P.O. Box 1539-UW2220
                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,110A
FILING DATE:
CONTRACTOR APPLICATION APPLICATION NUMBER: US/08/470,110A
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MEDIUM TYPE: Floppy disk
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COMPUTER READABLE FORM:
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                                                   CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                             71
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les 14; Conservative
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                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12 SGSGTDATAPASSLQAED 29
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                  19406-0939
                                               Pennsylvania
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                                                                                                                                                                                                                                                                                Ames, Robert S.
Appelbaum, Edward R.
Chaiken, Irwin M.
                                    USA
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                                                                                                                                                                                                             Lynette J.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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0.043;
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: LENGTH: 113 amino acids
: TYPE: amino acid
: STRANDEDNESS: single
: TOPOLOGY: linear
: MOLECULE TYPE: protein
US-08-667-769A-21
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US-08-940-371-21
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APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irvin M.
APPLICANT: Cook, Richard M.
APPLICANT: Cook, Richard M.
APPLICANT: Holmes, Stephen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21, Apprint No. 5851525
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
ZIP: 1940b-0000
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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FILING DATE: 06-7UN-195
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
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                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp./Corporate
ADDRESSEE: Intellectual Property
                                                                                                                                                                                                                                            APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy M.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
NUMBER OF SEQUENCES: 74
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CITY: King of Prussia
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                                                                                                     COUNTRY:
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Local Similarity 77.8%;
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                                                                                                                        Pennsylvania
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23-DEC-1994
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UMBER: 34,028
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Pred. No. 0.043;
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; MOLECULE TYPE: protein US-08-940-371-21
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Best Local :
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INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
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REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: APPLICANT:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
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               APPLICATION NUMBER: US/08/637,647
FILING DATE: 19-JUN-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: sir
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SOFTWARE: Patentin Release #1.0, Version #1.25
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ames, Robert S
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77.8%;
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Pred. No.
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0.043;
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                                                               US-08-525-539A-80
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Best Local Similarity
Query Match
Best Local Similarity
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                                                                                                                                        TELEFAX: (415) 494-079
TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/525,539A FILING DATE: 14-SEP-1995 CLASSIFICATION: 424 ATTORNEY/AGENT INFORMATION: NAME: DYLAN, TYLER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: /JJ
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: E
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: P5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
                                                                                                                                                                                                                                                                                                                                               COMPUTER: JBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: DO COUTO, FERNANDO APPLICANT: CERIANI, ROBERTO I APPLICANT: PETERSON, JERRY A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 113 amino
TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                            TELEPHONE: (415) V-
TELEPHONE: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: MUKKLOU.
CURRETT: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    71 SGSGTDFTLTISSLQAED 88
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                                                                                           TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                       REGISTRATION NUMBER:
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                                                                                                                            CENGTH:
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                                                                                                                            113 amino acids
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                                                                               linear
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77.8%;
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 43.8%;
77.8%;
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                                                                                                                                                                                                                                                     37,612
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   Score
Pred.
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   DB 4;
0.043;
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                  Length 113;
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RESULT 13
PCT-US93-08435-8
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                                                                                                       Sequence 8, Applicati
GENERAL INFORMATION:
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Best Local Similarity //...
14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (215) 540-581
INFORMATION FOR SEQ ID NO:
          APPLICANT:
APPLICANT:
TITLE OF IN
                                                        APPLICANT:
                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Howson and Howson
STREET: Box 457, 321 Norristown Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: U. S. Government, Secretary of
APPLICANT: the Army
TITLE OF INVENTION: Novel Antibodies for Conferring Passive
TITLE OF INVENTION: Immunity Against Infection by a Pathogen in Man
NUMBER OF SEQUENCES: 61
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: SBC P50107
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/941,654
FILING DATE: 09-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                     12 SGSGTDATAPASSLQAED 29
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STATE: PA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                        SGSGTDFTLTISSLQAED 88
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19477
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amino acid
          T: U. S. Government, Secretary of T: the Army INVENTION: Novel Antibodies for
                                                                                                                        Application PC/TUS9308435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             (215) 540-9200
(215) 540-801
                                                        U. S. Government, the Navy
                                                                      SmithKline Beecham, Corporation U. S. Government, Secretary of
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        Novel Antibodies for Conferring Passive
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Pred. No. 0.043;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                           TITLE OF INVENTION: Treatment NUMBER OF SEQUENCE.
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APPLICANT:
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APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: SB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 09-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
                                                                                                                                                                                    NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: protein
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PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                 APPLICANT:
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NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
                                                                                                                                    ADDRESSEE: SmithKline Beecham Corp./Corporate STREET: P.O. Box 1539-UW2220 CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 SGSGTDATAPASSLQAED 29
                                                                                        COUNTRY: USA
ZIP: 19406-0939
                                                                                                                        STATE:
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amino acid
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                                                                                                                     Pennsylvania
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Box 457, 321 Norrist
                                                                                                                                                                                                                                                                               Cook, Richard M. Gross, Mitchell S. Holmes, Stephen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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Chaiken, Irwin
                                                                                                                                                                                                                                                                                                                                                                 Ames, Robert S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43.8%;
77.8%;
                                                                                                                                                                                                                    Recombinant IL5 Antagonists Useful Treatment of IL5 Mediated Disorder.
                                                                                                                                                                                                                                                                                                                                                  Edward R.
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Gaps

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PRIOR APPLICATION DATA:

APPLICATION NUMBER: FILING DATE: CLASSIFICATION:

CURRENT APPLICATION DATA:

PCT/US95/17082A

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Best Local S
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
TYPE: amino acid
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FILING DATE: 06-JUN-1333
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
APPLICATION TATE: 23-DEC-1994
                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 500 kb
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordberfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Saiko HOSOKAWA
APPLICANT: TOShiaki TAGAWA
APPLICANT: Yoko HIRAKAWA
APPLICANT: NO. 5767246ihiko
APPLICANT: Kazuhiro NAGAIKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/470110 FILING DATE: 06-JUN-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESSE: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
             CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07
FILING DATE: June 29,
                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                         CITY: Washington STATE: D.C.
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APPLICATION NUMBER:
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14; Conservative
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5767246
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Specifically Binding to Surface Antigen of Cancer
Cell Membrane
42
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77.8%;
                                                                                             US/08/360,125
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0; Mismatches
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No. 0.043;
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; RELEVANT RESIDUES US-08-360-125-6
Search completed: July 15, 2002, Job time: 391 sec
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                                                                                                           Query Match
Best Local Similarity
Matches 14; Conserv
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LENGTH: 114 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warren M. Cheek, Jr. REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 202-371-8850
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLONE:
POSITION IN GENOME:
CHROMOSOME/SEGMENT:
                                                                                                                                                                                                                                                                                                                                                   PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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ORIGINAL SOURCE:
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TISSUE TYPE:
CELL TYPE: H
CELL TYPE: a
CELL LINE:
ORGANELLE:
                                                                                         12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                            DOCUMENT NUMBER:
FILING DATE:
PUBLICATION DATE:
                                                                                                                                                                                                                                                                                                                                                               IDENTIFICATION METHOD: OTHER INFORMATION:
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                                                                 71 SGSGTDFTLTISSLQAED 88
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                                                                                                                                                                                                                                                                    PAGES:
                                                                                                                                                                                                                                                                                               VOLUME:
                                                                                                                                                                                                                                                                                                           JOURNAL:
                                                                                                                                                                                                                                                                                                                                    AUTHORS:
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                                                                                                                                                                                                                                                                                  ESSUE:
                                                                                                                                                                                                                                                                                                                                                                                         LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INDIVIDUAL ISOLATE:
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                                                                                                                     Conservative
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linear
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77.8%;
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